

SEQUENZPROTOKOLL

<110> Biosyn Arzneimittel GmbH

<120> Nukleinsäuremolekül, umfassend eine für ein Hämocyanin kodierende Nukleinsäuresequenz

<130> PCT1153-01966

<140>

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<170> PatentIn Ver. 2.1

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<211> 1269

<212> DNA

<213> Haliotis tuberculata

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<213> Haliotis tuberculata

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2

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<213> *Haliotis tuberculata*

<400> 3

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<212> DNA

<213> *Haliotis tuberculata*

<400> 4

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3

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<213> *Haliotis tuberculata*

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<212> DNA

<213> *Haliotis tuberculata*

<400> 6

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4

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<212> DNA

<213> *Haliotis tuberculata*

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<213> *Haliotis tuberculata*

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5

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<211> 1003

<212> DNA

<213> *Haliotis tuberculata*

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<212> DNA

<213> *Haliotis tuberculata*

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6

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<212> DNA

<213> *Haliotis tuberculata*

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<210> 12

<211> 1255

<212> DNA

<213> *Haliotis tuberculata*

<400> 12

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tcacctgacg ggttccaagc cattgcctct tccatgctc tgccaccact ctgcccttca 180
ccatctgcag ctccaccgta tgettgtgt gtccacggca tggctacatt tcccagtg 240
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gagtttgag gagaaaacat tacaacagag agagaagtca ttgcagacaa actttttgtc 480
aaaggtggac aggtttttga taaactgggt cttcaaaca gccatcctag gcctgagcag 540
gaaaactact gtgactttga gattcagttt gaaattcttc acaacggcgt tcacacgtgg 600

```

7

```

gtcggaggca gtcgtaccta ctctatcgga catcttcatt acgcattcta cgaccctctt 660
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cagagagggc tctcgggtga tgaggctcac tgtgctctcg agcaaagag agaaccattg 780
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caccttcgat atgatgatga cttcacatc tctgtcagtc tgaccgcaa caacggaact 1200
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```

<210> 13

<211> 1248

<212> DNA

<213> *Haliotis tuberculata*

<400> 13

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gtgacataaa taccaggagc atgtcaccca accgtgttcg ccgtgagctg agcgatctgt 60
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ccaacggata ccaggctctt gcagccttcc atgggctacc agcaggctgc catgatagcc 180
ggggaaatga gatcgcatgt tgcattcacg ggatgccgac cttccccag tggcacagac 240
tgtacaccct gcagttggag atggctctga ggagacatgg atcatctgtc gccatcccct 300
actgggactg gacaaagcct atctccgaac tcccctcgct cttcaccagc cctgagtatt 360
atgacccatg gcatgatgct gtggtaaaca acccattctc caaaggtttt gtcaaatttg 420
caaataccta cacagtaaga gaccacagg agatgctgtt ccagctttgt gaacatggag 480
agtcaatcct ctatgagcaa actcttcttg ctcttgagca aaccgactac tgtgattttg 540
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```

<210> 14

<211> 1207

<212> DNA

<213> *Haliotis tuberculata*

<400> 14

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gcgagatgaa cgcccgcaag gttgcctgtt gtgctcacgg tatggcctcc ttcccacact 240
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8

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ctgtctggca agcactgcag aaataccgag gactcccata caacgaagca cactgtgaaa 720
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ccaagactaa tgccaggcct atcgattcat ttgattatga gaggtttaac tatcagtatg 840
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gagtcattga tcagctccat ctccagtatg attcagattt cagtttcagg gtgaagcttg 1140
ttgccaccaa tggcactgag ctttcatcag accttctcaa gtcaccaaca attgaacatg 1200
aacttg      1207

```

<210> 15

<211> 1546

<212> DNA

<213> *Haliotis tuberculata*

<400> 15

```

agccacaga ggaccagtgt aagaaacaga agtcactcgc caacatactg acggcaatgc 60
acactttcat gtaaggaag ttgattcgct gtccctggat gaagcaaaca acttgaagaa 120
tgccctttac aagctacaga acgaccacag tctaacggga tacgaagcaa tctctggtta 180
ccatggatac cccaatctgt gtccggaaga aggcgatgac aaaatacccc tgctgcgtcc 240
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tgagcacaat ggtgcactgc ttggtgttcc ttactgggac tggaaacaagg acctgtcgtc 360
actgccggcg ttcttctccg actccagcaa caacaatccc tacttcaagt accacatcgc 420
cgggtgttgt cacgacaccg tcagagagcc aactagtctt atatataacc agcccaaat 480
ccatggttat gattatctct attacctagc attgaccacg cttgaagaaa acaattactg 540
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ccacgtctat tctgtcaagc ctggtgacta ctatgttact ggaccacga gagaccttg 1500
ccagaatgca gatgtcagga ttcatatcca tgttgaggat gagtaa      1546

```

<210> 16

<211> 967

<212> DNA

<213> *Megathura crenulata*

<400> 16

```

ggcctaccgt actgggactg gactgaaccc atgacacaca ttccgggtct ggcaggaaac 60
aaaacttatg tggattctca tgggtgcatt cacacaaatc cttttcatag ttcagtgatt 120
gcatttgaag aaaatgctcc ccacacaaaa agacaaatag atcaaagact ctttaaacc 180
gctacctttg gacaccacac agacctgttc aaccagattt tgtatgcctt tgaacaagaa 240
gattactgtg actttgaagt ccaatttgag attaccata acacgattca cgcttgaca 300
ggaggaagcg aacatttctc aatgtcgtcc ctacattaca cagctttcga tcctttgttt 360
tactttcacc attctaacgt tgatcgtctt tgggccgttt ggcaagcctt acagatgaga 420

```

9

```

cggcataaac cctacagggc ccactgcgcc atatctctgg aacatatgca tctgaaacca 480
ttcgcccttt catctcccct taacaataac gaaaagactc atgccaatgc catgccaac 540
aagatctacg actatgaaaa tgtctcccat tacacatacg aagatttaac atttggaggc 600
atctctctgg aaaacataga aaagatgac cagcaaaacc agcaagaaga cagaatatat 660
gccggttttc tcttggtggt cctacgtact tcagcaaatg ttgatatctt cattaact 720
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ctcactgctg atggcgat ttcgaagttact gttgacatca ctgaagtcga tggaaactaaa 900
cttgcatcca gtcttattcc acatgcttct gtcattcgtg agcatgcacg tggtaagctg 960
aatagag

```

<210> 17

<211> 1242

<212> DNA

<213> Megathura crenulata

<400> 17

```

ttaaatttga caaagtgcc aggagtcgtc ttattcgaaa aaatgtagac cgtttgagcc 60
ccgaggagat gaatgaactt cgtaaagccc tagccttact gaaagaggac aaaagtgccg 120
gtggatttca gcagcttgggt gcattccatg gggagccaaa atggtgtcct agtcccgaag 180
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tcgacaaaac aacaacaaga agtggttcaga ataaactctt cgaacagcct gagtttggtc 420
attatacaag cattgccaaa caagtactgc tagcgttggga acaggacaat ttctgtgact 540
ttgaaatcca atatgagatt gcccataact acatccatgc acttgtagga ggcgctcagc 600
cttatgggtat ggcacgtctt cgtacactg cttttgatcc actattctac ttgcatcact 660
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acaacggttg taactgtgct gttacatcga tgagagaacc tttgcaacca tttggcctct 780
ctgccaatat caacacagac catgtaacca aggagcattc agtgccattc aacgtttttg 840
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acgatagagt cttcaaatat gacataaccg aaaaactcca cgatctaaag ctgcatgcag 1140
aagaccactt ctacattgac tatgaagtat ttgaccttaa accagcaagc ctgggaaaaag 1200
atgtgttcaa gcagccttca gtcattcatg aaccaagaat ag

```

<210> 18

<211> 1236

<212> DNA

<213> Megathura crenulata

<400> 18

```

gtcaccatga aggcgaagta tatcaagctg aagtaacttc tgccaaccgt attcgaaaaa 60
acattgaaaa tctgagcctt ggtgaactcg aaagtctgag agctgccttc ctggaaattg 120
aaaacgatgg aacttacgaa tcaatagcta aattccatgg tagccctggt ttgtgccagt 180
taaattggtaa ccccatctct tgtgtgtcc atggcatgcc aactttccct cactggcaca 240
gactgtacgt ggtgtgctgt gagaatgcc tctgaaaaaa aggatcatct gtagctgttc 300
cctattggga ctggacaaaa cgaatcgaac atttacctca cctgatttca gacgccactt 360
actacaattc caggcaacat cactatgaga caaaccatt ccatcatggc aaaatcacac 420
acgagaatga aatcactact agggatccca aggacagcct cttccattca gactactttt 480
acgagcaggt cttttacgcc ttggagcagg ataactctg tgatttcgag attcagttgg 540
agatattaca caatgcattg cattctttac ttggtggcaa aggtaaatat tccatgtcaa 600
accttgatta cgtgctttt gatcctgtgt tcttcttca tcacgcaacg actgacagaa 660
tctgggcaat ctggcaagac cttcagaggt tccgaaaacg gccataccga gaagcgaatt 720

```

10

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gcgctatcca attgatgcac acgccactcc agccggttga taagagcgac aacaatgacg 780
aggcaacgaa aacgcatgcc actccacatg atggttttga atatcaaaac agctttgggt 840
atgcttacga taatctggaa ctgaatcact actcgattcc tcagcttgat cacatgctgc 900
aagaaagaaa aaggcatgac agagtattcg ctggcttcct ccttcacaat attggaacat 960
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gactttacaa acttgacata actaaagcct tgaanaagaa cggtgtgcac ctgcaagggg 1140
atttcgatct ggaaattgag attacggctg tgaatggatc tcatctagac agtcatgtca 1200
tccactctcc cactatactg tttgaggccg gaacag 1236

```

<210> 19

<211> 241

<212> DNA

<213> *Megathura crenulata*

<400> 19

```

attctgcccc cacagatgat ggacacactg aaccagtgat gattcgcaaa gatatcacac 60
aattggacaa gcgtcaacaa ctgtcactgg tgaaagccct cgagtccatg aaagccgacc 120
attcatctga tgggttccag gcaatcgctt ccttccatgc tcttctctct ctttgtccat 180
caccagctgc ttcaaagagg tttgcgtgct gcgtccatgg catgccaaacc ttcccgaat 240
g 241

```

<210> 20

<211> 949

<212> DNA

<213> *Megathura crenulata*

<400> 20

```

ggcctgccct actgggattg gaccatgcca atgagtcatt tgccagaact ggctacaagt 60
gagacctacc tcgatccagt tactggggaa actaaaaaca accctttcca tcacgccccaa 120
gtggcggttg aaaatggtgt aacaagcagg aatcctgatg ccaaactttt tatgaaacca 180
acttacggag accacactta cctcttcgac agcatgatct acgcatttga gcaggaagac 240
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gcagtggatg atgggttcag cattactgtt gagatcaccg atgttgatgg atctccccc 900
tctgcagatc tcattccacc tctgctata atctttgaac gtggtcatg 949

```

<210> 21

<211> 760

<212> DNA

<213> *Megathura crenulata*

<400> 21

```

ctgatgccaa agactttggc catagcagaa aaatcaggaa agccgttgat tctctgacag 60
tcgaagaaca aacttcgttg aggcgagcta tggcagatct acaggacgac aaaacatcag 120
gggttttcca gcagattgca gcattccacg gagaacaaaa atggtgtcca agccccgaag 180
cggagaaaaa atttgcatgc tgtgttcatg gaatggctgt tttccctcac tggcacagat 240
tgctgacagt tcaaggagaa aatgctctga ggaaacatgg ctttactggg ggactgccct 300

```

11

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actgggactg gactcgatca atgagcgccc ttccacattt tgttgotgat cctacttaca 360
atgatgctat ttccagccag gaagaagata acccatggca tcatggtcac atagactctg 420
ttgggcatga tactacaaga gatgtgctg atgatcttta tcaatctcct ggtttcgggc 480
actacacaga tattgcacaa caagtccttc tggcctttga gcaggacagt ttctgtgatt 540
ttgaggtaca atttgaaatt gcccataaatt tcatacatgc actgattggg ggtaacgaac 600
catacagtat gtcactcttg aggtatacta catacgatec aatcttcttc ttgcaccact 660
ccagtacaga ccgacttttg gccatctggc aagcaatcac tagtgcggcc gcctgcaggt 720
cgaccataag ggagagctcc caacgcgttg gatgcaatct 760

```

<210> 22

<211> 323

<212> DNA

<213> Megathura crenulata

<400> 22

```

gttcacacca ggctgatgaa tatcgtgagg cagtaacaag cgctagccac ataagaaaaa 60
atatccggga cctctcagag ggagaaattg agagcatcag atctgctttc ctccaaattc 120
aaaaagaggg tatatatgaa aacattgcaa agttccatgg aaaaccagga ctttgtgaac 180
atgatggaca tcctgttgct tgttgtgtcc atggcatgcc cacctttccc cactggcaca 240
gactgtacgt tcttcaggtg gagaatgcgc tcttagaacg agggctctgca gttgctgttc 300
cttactggga ctggacccta cct 323

```

<210> 23

<211> 988

<212> DNA

<213> Megathura crenulata

<400> 23

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atggctgtgt ttccgcaactg gcacagactg tttgtgaaac agatggagga cgcacttgct 60
gctcatggag ctcatatttg cataccatac tgggattgga caagtgcgtt tagtcatctg 120
cccgccctag tgactgacca cgagaacaat cccttccacc acggccatat tggcatctg 180
aatgtggata catctcgatc tccaagagac atgctgttta atgatcctga acaaggctca 240
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gaagtgcagt ttgaacttac acacaatgcc atccactctt ggactggagg acatactcca 360
tatggaatgt catcactgga atatacagca tatgatccac tcttttatct ccaccattcc 420
aacactgatc gtatctgggc catctggcag gcaactccaga aatatagagg tcttccatac 480
aacgcagctc actgcgatat ccaagttctg aaacaacctc ttaaaccatt cagcgagtc 540
aggaatccaa acccagtcac cagagccaat tctagggccg ttgattcatt tgattatgag 600
aaattcaatt atcaatatga cacacttacc ttccacggac tttctatccc agaacttgat 660
gccatgcttc aagagagaaa gaaggaagag agaacatttg cagccttcct gttgcacgga 720
tttggcgcca gtgctgatgt ttcgtttgat gtctgcacac ctgatgggtca ttgtgccttt 780
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cactttgagt tgaagattgt tggcacagat ggaacagaac tgccatcgga tcgtatcaag 960
agccctacca ttgaacacca tggaggag 988

```

<210> 24

<211> 310

<212> DNA

<213> Megathura crenulata

<400> 24

```

gtcacgatca cagtgaacgt cacgatggat ttttcaggaa ggaagtcggt tccctgtccc 60
tggatgaagc caatgacctt aaaaatgcac tgtacaagct gcagaatgat caggggtccc 120
atggatatga atcaatagcc ggttaccatg gctatccatt cctctgccct gaacatgggt 180
aagaccagta cgcagctgtg gtccacggaa tgctgtatt tccacattgg cacagacttc 240
atacaatcca gtttgagaga gctctcaaag aacatgggtc tcatttgggt ctgccatact 300

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gggactggac

<210> 25

<211> 422

<212> PRT

<213> Haliotis tuberculata

<220>

<221> SIGNAL

<222> (1)..(15)

<400> 25

Leu Val Gln Phe Leu Leu Val Ala Leu Val Ala Gly Ala Gly Ala Asp
 1 5 10 15

Asn Val Val Arg Lys Asp Val Ser His Leu Thr Asp Asp Glu Val Gln
 20 25 30

Ala Leu His Gly Ala Leu His Asp Val Thr Ala Ser Thr Gly Pro Leu
 35 40 45

Ser Phe Glu Asp Ile Thr Ser Tyr His Ala Ala Pro Ala Ser Cys Asp
 50 55 60

Tyr Lys Gly Arg Lys Ile Ala Cys Cys Val His Gly Met Pro Ser Phe
 65 70 75 80

Pro Phe Trp His Arg Ala Tyr Val Val Gln Ala Glu Arg Ala Leu Leu
 85 90 95

Ser Lys Arg Lys Thr Val Gly Met Pro Tyr Trp Asp Trp Thr Gln Thr
 100 105 110

Leu Thr His Leu Pro Ser Leu Val Thr Glu Pro Ile Tyr Ile Asp Ser
 115 120 125

Lys Gly Gly Lys Ala Gln Thr Asn Tyr Trp Tyr Arg Gly Glu Ile Ala
 130 135 140

Phe Ile Asn Lys Lys Thr Ala Arg Ala Val Asp Asp Arg Leu Phe Glu
 145 150 155 160

Lys Val Glu Pro Gly His Tyr Thr His Leu Met Glu Thr Val Leu Asp
 165 170 175

Ala Leu Glu Gln Asp Glu Phe Cys Lys Phe Glu Ile Gln Phe Glu Leu
 180 185 190

Ala His Asn Ala Ile His Tyr Leu Val Gly Gly Lys Phe Glu Tyr Ser
 195 200 205

Met Ser Asn Leu Glu Tyr Thr Ser Tyr Asp Pro Ile Phe Phe Leu His
 210 215 220

His Ser Asn Val Asp Arg Leu Phe Ala Ile Trp Gln Arg Leu Gln Glu
 225 230 235 240

13

Leu Arg Gly Lys Asn Pro Asn Ala Met Asp Cys Ala His Glu Leu Ala
 245 250 255
 His Gln Gln Leu Gln Pro Phe Asn Arg Asp Ser Asn Pro Val Gln Leu
 260 265 270
 Thr Lys Asp His Ser Thr Pro Ala Asp Leu Phe Asp Tyr Lys Gln Leu
 275 280 285
 Gly Tyr Ser Tyr Asp Ser Leu Asn Leu Asn Gly Met Thr Pro Glu Gln
 290 295 300
 Leu Lys Thr Glu Leu Asp Glu Arg His Ser Lys Glu Arg Ala Phe Ala
 305 310 315 320
 Ser Phe Arg Leu Ser Gly Phe Gly Gly Ser Ala Asn Val Val Val Tyr
 325 330 335
 Ala Cys Val Pro Asp Asp Asp Pro Arg Ser Asp Asp Tyr Cys Glu Lys
 340 345 350
 Ala Gly Asp Phe Phe Ile Leu Gly Gly Gln Ser Glu Met Pro Trp Arg
 355 360 365
 Phe Tyr Arg Pro Phe Phe Tyr Asp Val Thr Glu Ala Val His His Leu
 370 375 380
 Gly Val Pro Leu Ser Gly His Tyr Tyr Val Lys Thr Glu Leu Phe Ser
 385 390 395 400
 Val Asn Gly Thr Ala Leu Ser Pro Asp Leu Leu Pro Gln Pro Thr Val
 405 410 415
 Ala Tyr Arg Pro Gly Lys
 420

<210> 26

<211> 419

<212> PRT

<213> *Haliotis tuberculata*

<400> 26

Gly His Leu Asp Pro Pro Val His His Arg His Asp Asp Asp Leu Ile
 1 5 10 15
 Val Arg Lys Asn Ile Asp His Leu Thr Arg Glu Glu Glu Tyr Glu Leu
 20 25 30
 Arg Met Ala Leu Glu Arg Phe Gln Ala Asp Thr Ser Val Asp Gly Tyr
 35 40 45
 Gln Ala Thr Val Glu Tyr His Gly Leu Pro Ala Arg Cys Pro Arg Pro
 50 55 60
 Asp Ala Lys Val Arg Phe Ala Cys Cys Met His Gly Met Ala Ser Phe
 65 70 75 80

Pro His Trp His Arg Leu Phe Val Thr Gln Val Glu Asp Ala Leu Val
85 90 95

Arg Arg Gly Ser Pro Ile Gly Val Pro Tyr Trp Asp Trp Thr Lys Pro
100 105 110

Met Thr His Leu Pro Asp Leu Ala Ser Asn Glu Thr Tyr Val Asp Pro
115 120 125

Tyr Gly His Thr His His Asn Pro Phe Phe Asn Ala Asn Ile Ser Phe
130 135 140

Glu Glu Gly His His His Thr Ser Arg Met Ile Asp Ser Lys Leu Phe
145 150 155 160

Ala Pro Val Ala Phe Gly Glu His Ser His Leu Phe Asp Gly Ile Leu
165 170 175

Tyr Ala Phe Glu Gln Glu Asp Phe Cys Asp Phe Glu Ile Gln Phe Glu
180 185 190

Leu Val His Asn Ser Ile His Ala Trp Ile Gly Gly Ser Glu Asp Tyr
195 200 205

Ser Met Ala Thr Leu His Tyr Thr Ala Phe Asp Pro Ile Phe Tyr Leu
210 215 220

His His Ser Asn Val Asp Arg Leu Trp Ala Ile Trp Gln Ala Leu Gln
225 230 235 240

Ile Arg Arg His Lys Pro Tyr Gln Ala His Cys Ala Gln Ser Val Glu
245 250 255

Gln Leu Pro Met Lys Pro Phe Ala Phe Pro Ser Pro Leu Asn Asn Asn
260 265 270

Glu Lys Thr His Ser His Ser Val Pro Thr Asp Ile Tyr Asp Tyr Glu
275 280 285

Glu Val Leu His Tyr Ser Tyr Asp Asp Leu Thr Phe Gly Gly Met Asn
290 295 300

Leu Glu Glu Ile Glu Glu Ala Ile His Leu Arg Gln Gln His Glu Arg
305 310 315 320

Val Phe Ala Gly Phe Leu Leu Ala Gly Ile Gly Thr Ser Ala Leu Val
325 330 335

Asp Ile Phe Ile Asn Lys Pro Gly Asn Gln Pro Leu Lys Ala Gly Asp
340 345 350

Ile Ala Ile Leu Gly Gly Ala Lys Glu Met Pro Trp Ala Phe Asp Arg
355 360 365

Leu Tyr Lys Val Glu Ile Thr Asp Ser Leu Lys Thr Leu Ser Leu Asp
370 375 380

15

Val Asp Gly Asp Tyr Glu Val Thr Phe Lys Ile His Asp Met His Gly
385 390 395 400

Asn Ala Leu Asp Thr Asp Leu Ile Pro His Ala Ala Val Val Ser Glu
405 410 415

Pro Ala His

<210> 27

<211> 414

<212> PRT

<213> *Haliotis tuberculata*

<400> 27

Pro Thr Phe Glu Asp Glu Lys His Ser Leu Arg Ile Arg Lys Asn Val
1 5 10 15

Asp Ser Leu Thr Pro Glu Glu Thr Asn Glu Leu Arg Lys Ala Leu Glu
20 25 30

Leu Leu Glu Asn Asp His Thr Ala Gly Gly Phe Asn Gln Leu Gly Ala
35 40 45

Phe His Gly Glu Pro Lys Trp Cys Pro Asn Pro Glu Ala Glu His Lys
50 55 60

Val Ala Cys Cys Val His Gly Met Ala Val Phe Pro His Trp His Arg
65 70 75 80

Leu Leu Ala Leu Gln Ala Glu Asn Ala Leu Arg Lys His Gly Tyr Ser
85 90 95

Gly Ala Leu Pro Tyr Trp Asp Trp Thr Arg Pro Leu Ser Gln Leu Pro
100 105 110

Asp Leu Val Ser His Glu Gln Tyr Thr Asp Pro Ser Asp His His Val
115 120 125

Lys His Asn Pro Trp Phe Asn Gly His Ile Asp Thr Val Asn Gln Asp
130 135 140

Thr Thr Arg Ser Val Arg Glu Asp Leu Tyr Gln Gln Pro Glu Phe Gly
145 150 155 160

His Phe Thr Asp Ile Ala Gln Gln Val Leu Leu Ala Leu Glu Gln Asp
165 170 175

Asp Phe Cys Ser Phe Glu Val Gln Tyr Glu Ile Ser His Asn Phe Ile
180 185 190

His Ala Leu Val Gly Gly Thr Asp Ala Tyr Gly Met Ala Ser Leu Arg
195 200 205

16

Tyr Thr Ala Tyr Asp Pro Ile Phe Phe Leu His His Ser Asn Thr Asp
 210 215 220
 Arg Ile Trp Ala Ile Trp Gln Ser Leu Gln Lys Tyr Arg Gly Lys Pro
 225 230 235 240
 Tyr Asn Thr Ala Asn Cys Ala Ile Glu Ser Met Arg Arg Pro Leu Gln
 245 250 255
 Pro Phe Gly Leu Ser Ser Ala Ile Asn Pro Asp Arg Ile Thr Arg Glu
 260 265 270
 His Ala Ile Pro Phe Asp Val Phe Asn Tyr Arg Asp Asn Leu His Tyr
 275 280 285
 Val Tyr Asp Thr Leu Glu Phe Asn Gly Leu Ser Ile Ser Gln Leu Asp
 290 295 300
 Arg Glu Leu Glu Lys Ile Lys Ser His Glu Arg Val Phe Ala Gly Phe
 305 310 315 320
 Leu Leu Ser Gly Ile Lys Lys Ser Ala Leu Val Lys Phe Glu Val Cys
 325 330 335
 Thr Pro Pro Asp Asn Cys His Lys Ala Gly Glu Phe Tyr Leu Leu Gly
 340 345 350
 Asp Glu Asn Glu Met Ala Trp Ala Tyr Asp Arg Leu Phe Lys Tyr Asp
 355 360 365
 Ile Thr Gln Val Leu Glu Ala Asn His Leu His Phe Tyr Asp His Leu
 370 375 380
 Phe Ile Arg Tyr Glu Val Phe Asp Leu Lys Gly Val Ser Leu Gly Thr
 385 390 395 400
 Asp Leu Phe His Thr Ala Asn Val Val His Asp Ser Gly Thr
 405 410

<210> 28

<211> 413

<212> PRT

<213> Haliotis tuberculata

<400> 28

Gly Thr Arg Asp Arg Asp Asn Tyr Val Glu Glu Val Thr Gly Ala Ser
 1 5 10 15
 His Ile Arg Lys Asn Leu Asn Asp Leu Asn Thr Gly Glu Met Glu Ser
 20 25 30
 Leu Arg Ala Ala Phe Leu His Ile Gln Asp Asp Gly Thr Tyr Glu Ser
 35 40 45
 Ile Ala Gln Tyr His Gly Lys Pro Gly Lys Cys Gln Leu Asn Asp His
 50 55 60

17

Asn Ile Ala Cys Cys Val His Gly Met Pro Thr Phe Pro Gln Trp His
 65 70 75 80
 Arg Leu Tyr Val Val Gln Val Glu Asn Ala Leu Leu Asn Arg Gly Ser
 85 90 95
 Gly Val Ala Val Pro Tyr Trp Glu Trp Thr Ala Pro Ile Asp His Leu
 100 105 110
 Pro His Phe Ile Asp Asp Ala Thr Tyr Phe Asn Ser Arg Gln Gln Arg
 115 120 125
 Tyr Asp Pro Asn Pro Phe Phe Arg Gly Lys Val Thr Phe Glu Asn Ala
 130 135 140
 Val Thr Thr Arg Asp Pro Gln Ala Gly Leu Phe Asn Ser Asp Tyr Met
 145 150 155 160
 Tyr Glu Asn Val Leu Leu Ala Leu Glu Gln Glu Asn Tyr Cys Asp Phe
 165 170 175
 Glu Ile Gln Phe Glu Leu Val His Asn Ala Leu His Ser Met Leu Gly
 180 185 190
 Gly Lys Gly Gln Tyr Ser Met Ser Ser Leu Asp Tyr Ser Ala Phe Asp
 195 200 205
 Pro Val Phe Phe Leu His His Ala Asn Thr Asp Arg Leu Trp Ala Ile
 210 215 220
 Trp Gln Glu Leu Gln Arg Phe Arg Glu Leu Pro Tyr Glu Glu Ala Asn
 225 230 235 240
 Cys Ala Ile Asn Leu Met His Gln Pro Leu Lys Pro Phe Ser Asp Pro
 245 250 255
 His Glu Asn His Asp Asn Val Thr Leu Lys Tyr Ser Lys Pro Gln Asp
 260 265 270
 Gly Phe Asp Tyr Gln Asn His Phe Gly Tyr Lys Tyr Asp Asn Leu Glu
 275 280 285
 Phe His His Leu Ser Ile Pro Ser Leu Asp Ala Thr Leu Lys Gln Arg
 290 295 300
 Arg Asn His Asp Arg Val Phe Ala Gly Phe Leu Leu His Asn Ile Gly
 305 310 315 320
 Thr Ser Ala Asp Ile Thr Ile Tyr Ile Cys Leu Pro Asp Gly Arg Arg
 325 330 335
 Gly Asn Asp Cys Ser His Glu Ala Gly Thr Phe Tyr Ile Leu Gly Gly
 340 345 350
 Glu Thr Glu Met Pro Phe Ile Phe Asp Arg Leu Tyr Lys Phe Glu Ile
 355 360 365

Thr Lys Pro Leu Gln Gln Leu Gly Val Lys Leu His Gly Gly Val Phe
 370 375 380

Glu Leu Glu Leu Glu Ile Lys Ala Tyr Asn Gly Ser Tyr Leu Asp Pro
 385 390 395 400

His Thr Phe Asp Pro Thr Ile Ile Phe Glu Pro Gly Thr
 405 410

<210> 29

<211> 420

<212> PRT

<213> *Haliotis tuberculata*

<400> 29

Asp Thr His Ile Leu Asp His Asp His Glu Glu Glu Ile Leu Val Arg
 1 5 10 15

Lys Asn Ile Ile Asp Leu Ser Pro Arg Glu Arg Val Ser Leu Val Lys
 20 25 30

Ala Leu Gln Arg Met Lys Asn Asp Arg Ser Ala Asp Gly Tyr Gln Ala
 35 40 45

Ile Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Asn Pro Ser Ala
 50 55 60

Ala His Arg Tyr Ala Cys Cys Val His Gly Met Ala Thr Phe Pro Gln
 65 70 75 80

Trp His Arg Leu Tyr Thr Val Gln Val Gln Asp Ala Leu Arg Arg His
 85 90 95

Gly Ser Leu Val Gly Ile Pro Tyr Trp Asp Trp Thr Lys Pro Val Asn
 100 105 110

Glu Leu Pro Glu Leu Leu Ser Ser Ala Thr Phe Tyr His Pro Ile Arg
 115 120 125

Asn Ile Asn Ile Ser Asn Pro Phe Leu Gly Ala Asp Ile Glu Phe Glu
 130 135 140

Gly Pro Gly Val His Thr Glu Arg His Ile Asn Thr Glu Arg Leu Phe
 145 150 155 160

His Ser Gly Asp His Asp Gly Tyr His Asn Trp Phe Phe Glu Thr Val
 165 170 175

Leu Phe Ala Leu Glu Gln Glu Asp Tyr Cys Asp Phe Glu Ile Gln Phe
 180 185 190

Glu Ile Ala His Asn Gly Ile His Thr Trp Ile Gly Gly Ser Ala Val
 195 200 205

19

Tyr Gly Met Gly His Leu His Tyr Ala Ser Tyr Asp Pro Ile Phe Tyr
 210 215 220
 Ile His His Ser Gln Thr Asp Arg Ile Trp Ala Ile Trp Gln Glu Leu
 225 230 235 240
 Gln Lys Tyr Arg Gly Leu Ser Gly Ser Glu Ala Asn Cys Ala Ile Glu
 245 250 255
 His Met Arg Thr Pro Leu Lys Pro Phe Ser Phe Gly Pro Pro Tyr Asn
 260 265 270
 Leu Asn Ser His Thr Gln Glu Tyr Ser Lys Pro Glu Asp Thr Phe Asp
 275 280 285
 Tyr Lys Lys Phe Gly Tyr Arg Tyr Asp Ser Leu Glu Leu Glu Gly Arg
 290 295 300
 Ser Ile Ser Arg Ile Asp Glu Leu Ile Gln Gln Arg Gln Glu Lys Asp
 305 310 315 320
 Arg Thr Phe Ala Gly Phe Leu Leu Lys Gly Phe Gly Thr Ser Ala Ser
 325 330 335
 Val Ser Leu Gln Val Cys Arg Val Asp His Thr Cys Lys Asp Ala Gly
 340 345 350
 Tyr Phe Thr Ile Leu Gly Gly Ser Ala Glu Met Pro Trp Ala Phe Asp
 355 360 365
 Arg Leu Tyr Lys Tyr Asp Ile Thr Lys Thr Leu His Asp Met Asn Leu
 370 375 380
 Arg His Glu Asp Thr Phe Ser Ile Asp Val Thr Ile Thr Ser Tyr Asn
 385 390 395 400
 Gly Thr Val Leu Ser Gly Asp Leu Ile Gln Thr Pro Ser Ile Ile Phe
 405 410 415
 Val Pro Gly Arg
 420

<210> 30

<211> 417

<212> PRT

<213> Haliotis tuberculata

<400> 30

His Lys Leu Asn Ser Arg Lys His Thr Pro Asn Arg Val Arg His Glu
 1 5 10 15

Leu Ser Ser Leu Ser Ser Arg Asp Ile Ala Ser Leu Lys Ala Ala Leu
 20 25 30

Thr Ser Leu Gln His Asp Asn Gly Thr Asp Gly Tyr Gln Ala Ile Ala
 35 40 45

20

Ala Phe His Gly Val Pro Ala Gln Cys His Glu Pro Ser Gly Arg Glu
 50 55 60
 Ile Ala Cys Cys Ile His Gly Met Ala Thr Phe Pro His Trp His Arg
 65 70 75 80
 Leu Tyr Thr Leu Gln Leu Glu Gln Ala Leu Arg Arg His Gly Ser Ser
 85 90 95
 Val Ala Val Pro Tyr Trp Asp Trp Thr Lys Pro Ile Thr Glu Leu Pro
 100 105 110
 His Ile Leu Thr Asp Gly Glu Tyr Tyr Asp Val Trp Gln Asn Ala Val
 115 120 125
 Leu Ala Asn Pro Phe Ala Arg Gly Tyr Val Lys Ile Lys Asp Ala Phe
 130 135 140
 Thr Val Arg Asn Val Gln Glu Ser Leu Phe Lys Met Ser Ser Phe Gly
 145 150 155 160
 Lys His Ser Leu Leu Phe Asp Gln Ala Leu Leu Ala Leu Glu Gln Thr
 165 170 175
 Asp Tyr Cys Asp Phe Glu Val Gln Phe Glu Val Met His Asn Thr Ile
 180 185 190
 His Tyr Leu Val Gly Gly Arg Gln Thr Tyr Ala Phe Ser Ser Leu Glu
 195 200 205
 Tyr Ser Ser Tyr Asp Pro Ile Phe Phe Ile His His Ser Phe Val Asp
 210 215 220
 Lys Ile Trp Ala Val Trp Gln Glu Leu Gln Ser Arg Arg His Leu Gln
 225 230 235 240
 Phe Arg Thr Ala Asp Cys Ala Val Gly Leu Met Gly Gln Ala Met Arg
 245 250 255
 Pro Phe Asn Lys Asp Phe Asn His Asn Ser Phe Thr Lys Lys His Ala
 260 265 270
 Val Pro Asn Thr Val Phe Asp Tyr Glu Asp Leu Gly Tyr Asn Tyr Asp
 275 280 285
 Asn Leu Glu Ile Ser Gly Leu Asn Leu Asn Glu Ile Glu Ala Leu Ile
 290 295 300
 Ala Lys Arg Lys Ser His Ala Arg Val Phe Ala Gly Phe Leu Leu Phe
 305 310 315 320
 Gly Leu Gly Thr Ser Ala Asp Ile His Leu Glu Ile Cys Lys Thr Ser
 325 330 335
 Glu Asn Cys His Asp Ala Gly Val Ile Phe Ile Leu Gly Gly Ser Ala
 340 345 350

ERSATZBLATT (REGEL 26)

21

Glu Met His Trp Ala Tyr Asn Arg Leu Tyr Lys Tyr Asp Ile Thr Glu
 355 360 365
 Ala Leu Gln Glu Phe Asp Ile Asn Pro Glu Asp Val Phe His Ala Asp
 370 375 380
 Glu Pro Phe Phe Leu Arg Leu Ser Val Val Ala Val Asn Gly Thr Val
 385 390 395 400
 Ile Pro Ser Ser His Leu His Gln Pro Thr Ile Ile Tyr Glu Pro Gly
 405 410 415
 Glu

<210> 31
 <211> 403
 <212> PRT
 <213> Haliotis tuberculata

<400> 31
 Asp His His Asp Asp His Gln Ser Gly Ser Ile Ala Gly Ser Gly Val
 1 5 10 15
 Arg Lys Asp Val Asn Thr Leu Thr Lys Ala Glu Thr Asp Asn Leu Arg
 20 25 30
 Glu Ala Leu Trp Gly Val Met Ala Asp His Gly Pro Asn Gly Phe Gln
 35 40 45
 Ala Ile Ala Ala Phe His Gly Lys Pro Ala Leu Cys Pro Met Pro Asp
 50 55 60
 Gly His Asn Tyr Ser Cys Cys Thr His Gly Met Ala Thr Phe Pro His
 65 70 75 80
 Trp His Arg Leu Tyr Thr Lys Gln Met Glu Asp Ala Met Arg Ala His
 85 90 95
 Gly Ser His Val Gly Leu Pro Tyr Trp Asp Trp Thr Ala Ala Phe Thr
 100 105 110
 His Leu Pro Thr Leu Val Thr Asp Thr Asp Asn Asn Pro Phe Gln His
 115 120 125
 Gly His Ile Asp Tyr Leu Asn Val Ser Thr Thr Arg Ser Pro Arg Asp
 130 135 140
 Met Leu Phe Asn Asp Pro Glu His Gly Ser Glu Ser Phe Phe Tyr Arg
 145 150 155 160
 Gln Val Leu Leu Ala Leu Glu Gln Thr Asp Phe Cys Lys Phe Glu Val
 165 170 175

22

Gln Phe Glu Ile Thr His Asn Ala Ile His Ser Trp Thr Gly Gly His
 180 185 190
 Ser Pro Tyr Gly Met Ser Thr Leu Asp Phe Thr Ala Tyr Asp Pro Leu
 195 200 205
 Phe Trp Leu His His Ser Asn Thr Asp Arg Ile Trp Ala Val Trp Gln
 210 215 220
 Ala Leu Gln Glu Tyr Arg Gly Leu Pro Tyr Asn His Ala Asn Cys Glu
 225 230 235 240
 Ile Gln Ala Met Lys Thr Pro Leu Arg Pro Phe Ser Asp Asp Ile Asn
 245 250 255
 His Asn Pro Val Thr Lys Ala Asn Ala Lys Pro Leu Asp Val Phe Glu
 260 265 270
 Tyr Asn Arg Leu Ser Phe Gln Tyr Asp Asn Leu Ile Phe His Gly Tyr
 275 280 285
 Ser Ile Pro Glu Leu Asp Arg Val Leu Glu Glu Arg Lys Glu Glu Asp
 290 295 300
 Arg Ile Phe Ala Ala Phe Leu Leu Ser Gly Ile Lys Arg Ser Ala Asp
 305 310 315 320
 Val Val Phe Asp Ile Cys Gln Pro Glu His Glu Cys Val Phe Ala Gly
 325 330 335
 Thr Phe Ala Ile Leu Gly Gly Glu Leu Glu Met Pro Trp Ser Phe Asp
 340 345 350
 Arg Leu Phe Arg Tyr Asp Ile Thr Lys Val Met Lys Gln Leu His Leu
 355 360 365
 Arg His Asp Ser Asp Phe Thr Phe Arg Val Lys Ile Val Gly Thr Asp
 370 375 380
 Asp His Glu Leu Pro Ser Asp Ser Val Lys Ala Pro Thr Ile Glu Phe
 385 390 395 400
 Glu Pro Gly

<210> 32

<211> 511

<212> PRT

<213> Haliotis tuberculata

<400> 32

Val His Arg Gly Gly Asn His Glu Asp Glu His His Asp Asp Arg Leu
 1 5 10 15

Ala Asp Val Leu Ile Arg Lys Glu Val Asp Phe Leu Ser Leu Gln Glu
 20 25 30

23

Ala Asn Ala Ile Lys Asp Ala Leu Tyr Lys Leu Gln Asn Asp Asp Ser
 35 40 45
 Lys Gly Gly Phe Glu Ala Ile Ala Gly Tyr His Gly Tyr Pro Asn Met
 50 55 60
 Cys Pro Glu Arg Gly Thr Asp Lys Tyr Pro Cys Cys Val His Gly Met
 65 70 75 80
 Pro Val Phe Pro His Trp His Arg Leu His Thr Ile Gln Met Glu Arg
 85 90 95
 Ala Leu Lys Asn His Gly Ser Pro Met Gly Ile Pro Tyr Trp Asp Trp
 100 105 110
 Thr Lys Lys Met Ser Ser Leu Pro Ser Phe Phe Gly Asp Ser Ser Asn
 115 120 125
 Asn Asn Pro Phe Tyr Lys Tyr Tyr Ile Arg Gly Val Gln His Glu Thr
 130 135 140
 Thr Arg Asp Val Asn Gln Arg Leu Phe Asn Gln Thr Lys Phe Gly Glu
 145 150 155 160
 Phe Asp Tyr Leu Tyr Tyr Leu Thr Leu Gln Val Leu Glu Glu Asn Ser
 165 170 175
 Tyr Cys Asp Phe Glu Val Gln Tyr Glu Ile Leu His Asn Ala Val His
 180 185 190
 Ser Trp Leu Gly Gly Thr Gly Gln Tyr Ser Met Ser Thr Leu Glu Tyr
 195 200 205
 Ser Ala Phe Asp Pro Val Phe Met Ile His His Ser Ser Leu Asp Arg
 210 215 220
 Ile Trp Ile Leu Trp Gln Lys Leu Gln Lys Ile Arg Met Lys Pro Tyr
 225 230 235 240
 Tyr Ala Leu Asp Cys Ala Gly Asp Arg Leu Met Lys Asp Pro Leu His
 245 250 255
 Pro Phe Asn Tyr Glu Thr Val Asn Glu Asp Glu Phe Thr Arg Ile Asn
 260 265 270
 Ser Phe Pro Ser Ile Leu Phe Asp His Tyr Arg Phe Asn Tyr Glu Tyr
 275 280 285
 Asp Asn Met Arg Ile Arg Gly Gln Asp Ile His Glu Leu Glu Glu Val
 290 295 300
 Ile Gln Glu Leu Arg Asn Lys Asp Arg Ile Phe Ala Gly Phe Val Leu
 305 310 315 320
 Ser Gly Leu Arg Ile Ser Ala Thr Val Lys Val Phe Ile His Ser Lys
 325 330 335

24

Asn Asp Thr Ser His Glu Glu Tyr Ala Gly Glu Phe Ala Val Leu Gly
 340 345 350
 Gly Glu Lys Glu Met Pro Trp Ala Tyr Glu Arg Met Leu Lys Leu Asp
 355 360 365
 Ile Ser Asp Ala Val His Lys Leu His Val Lys Asp Glu Asp Ile Arg
 370 375 380
 Phe Arg Val Val Val Thr Ala Tyr Asn Gly Asp Val Val Thr Thr Arg
 385 390 395 400
 Leu Ser Gln Pro Phe Ile Val His Arg Pro Ala His Val Ala His Asp
 405 410 415
 Ile Leu Val Ile Pro Val Gly Ala Gly His Asp Leu Pro Pro Lys Val
 420 425 430
 Val Val Lys Ser Gly Thr Lys Val Glu Phe Thr Pro Ile Asp Ser Ser
 435 440 445
 Val Asn Lys Ala Met Val Glu Leu Gly Ser Tyr Thr Ala Met Ala Lys
 450 455 460
 Cys Ile Val Pro Pro Phe Ser Tyr His Gly Phe Glu Leu Asp Lys Val
 465 470 475 480
 Tyr Ser Val Asp His Gly Asp Tyr Tyr Ile Ala Ala Gly Thr His Ala
 485 490 495
 Leu Cys Glu Gln Asn Leu Arg Leu His Ile His Val Glu His Glu
 500 505 510

<210> 33

<211> 334

<212> PRT

<213> Haliotis tuberculata

<400> 33

His Arg Leu Phe Val Thr Gln Val Glu Asp Ala Leu Ile Arg Arg Gly
 1 5 10 15
 Ser Pro Ile Gly Val Pro Tyr Trp Asp Trp Thr Gln Pro Met Ala His
 20 25 30
 Leu Pro Gly Leu Ala Asp Asn Ala Thr Tyr Arg Asp Pro Ile Ser Gly
 35 40 45
 Asp Ser Arg His Asn Pro Phe His Asp Val Glu Val Ala Phe Glu Asn
 50 55 60
 Gly Arg Thr Glu Arg His Pro Asp Ser Arg Leu Phe Glu Gln Pro Leu
 65 70 75 80

Phe	Gly	Lys	His	Thr	Arg	Leu	Phe	Asp	Ser	Ile	Val	Tyr	Ala	Phe	Glu
85								90				95			
Gln	Glu	Asp	Phe	Cys	Asp	Phe	Glu	Val	Gln	Phe	Glu	Met	Thr	His	Asn
100								105				110			
Asn	Ile	His	Ala	Trp	Ile	Gly	Gly	Gly	Glu	Lys	Tyr	Ser	Met	Ser	Ser
115								120				125			
Leu	His	Tyr	Thr	Ala	Phe	Asp	Pro	Ile	Phe	Tyr	Leu	Arg	His	Ser	Asn
130								135				140			
Thr	Asp	Arg	Leu	Trp	Ala	Ile	Trp	Gln	Ala	Leu	Gln	Ile	Arg	Arg	Asn
145								150				155			
Arg	Pro	Tyr	Lys	Ala	His	Cys	Ala	Trp	Ser	Glu	Glu	Arg	Gln	Pro	Leu
				165				170				175			
Lys	Pro	Phe	Ala	Phe	Ser	Ser	Pro	Leu	Asn	Asn	Asn	Glu	Lys	Thr	Tyr
				180				185				190			
Glu	Asn	Ser	Val	Pro	Thr	Asn	Val	Tyr	Asp	Tyr	Glu	Gly	Val	Leu	Gly
				195				200				205			
Tyr	Thr	Tyr	Asp	Asp	Leu	Asn	Phe	Gly	Gly	Met	Asp	Leu	Gly	Gln	Leu
210				215				220							
Glu	Glu	Tyr	Ile	Gln	Arg	Gln	Arg	Gln	Arg	Asp	Arg	Thr	Phe	Ala	Gly
225				230				235				240			
Phe	Phe	Leu	Ser	His	Ile	Gly	Thr	Ser	Ala	Asn	Val	Glu	Ile	Ile	Ile
				245				250				255			
Asp	His	Gly	Thr	Leu	His	Thr	Ser	Val	Gly	Thr	Phe	Ala	Val	Leu	Gly
				260				265				270			
Gly	Glu	Lys	Glu	Met	Lys	Trp	Gly	Phe	Asp	Arg	Leu	Tyr	Lys	Tyr	Glu
				275				280				285			
Ile	Thr	Asp	Glu	Leu	Arg	Gln	Leu	Asn	Leu	Arg	Ala	Asp	Asp	Val	Phe
290				295				300							
Ser	Ile	Ser	Val	Lys	Val	Thr	Asp	Val	Asp	Gly	Ser	Glu	Leu	Ser	Ser
305				310				315				320			
Glu	Leu	Ile	Pro	Ser	Ala	Ala	Ile	Ile	Phe	Glu	Arg	Ser	His		
				325				330							

<400> 34
Ile Asp His Gln Asp Pro His His Asp Thr Ile Ile Arg Lys Asn Val
1 5 10 15

26

Asp Asn Leu Thr Pro Glu Glu Ile Asn Ser Leu Arg Arg Ala Met Ala
 20 25 30
 Asp Leu Gln Ser Asp Lys Thr Ala Gly Gly Phe Gln Gln Ile Ala Ala
 35 40 45
 Phe His Gly Glu Pro Lys Trp Cys Pro Ser Pro Asp Ala Glu Lys Lys
 50 55 60
 Phe Ser Cys Cys Val His Gly Met Ala Val Phe Pro His Trp His Arg
 65 70 75 80
 Leu Leu Thr Val Gln Gly Glu Asn Ala Leu Arg Lys His Gly Cys Leu
 85 90 95
 Gly Ala Leu Pro Tyr Trp Asp Trp Thr Arg Pro Leu Ser His Leu Pro
 100 105 110
 Asp Leu Val Leu Val Ser Ser Arg Thr Thr Pro Met Pro Tyr Ser Thr
 115 120 125
 Val Glu Ala Arg Asn Pro Trp Tyr Ser Gly His Ile Asp Thr Val Gly
 130 135 140
 Val Asp Thr Thr Arg Ser Val Arg Gln Glu Leu Tyr Glu Ala Pro Gly
 145 150 155 160
 Phe Gly His Tyr Thr Gly Val Ala Lys Gln Val Leu Leu Ala Leu Glu
 165 170 175
 Gln Asp Asp Phe Cys Asp Phe Glu Val Gln Phe Glu Ile Ala His Asn
 180 185 190
 Phe Ile His Ala Leu Val Gly Gly Ser Glu Pro Tyr Gly Met Ala Ser
 195 200 205
 Leu Arg Tyr Thr Thr Tyr Asp Pro Ile Phe Tyr Leu His His Ser Asn
 210 215 220
 Thr Asp Arg Leu Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly
 225 230 235 240
 Lys Pro Tyr Asn Ser Ala Asn Cys Ala Ile Ala Ser Met Arg Lys Pro
 245 250 255
 Leu Gln Pro Phe Gly Leu Thr Asp Glu Ile Asn Pro Asp Asp Glu Thr
 260 265 270
 Arg Gln His Ala Val Pro Phe Ser Val Phe Asp Tyr Lys Asn Asn Phe
 275 280 285
 Asn Tyr Glu Tyr Asp Thr Leu Asp Phe Asn Gly Leu Ser Ile Ser Gln
 290 295 300
 Leu Asp Arg Glu Leu Ser Arg Arg Lys Ser His Asp Arg Val Phe Ala
 305 310 315 320

ERSATZBLATT (REGEL 26)

27

Gly Phe Leu Leu His Gly Ile Gln Gln Ser Ala Leu Val Lys Phe Phe
325 330 335

Val Cys Lys Ser Asp Asp Asp Cys Asp His Tyr Ala Gly Glu Phe Tyr
340 345 350

Ile Leu Gly Asp Glu Ala Glu Met Pro Trp Gly Tyr Asp Arg Leu Tyr
355 360 365

Lys Tyr Glu Ile Thr Glu Gln Leu Asn Ala Leu Asp Leu His Ile Gly
370 375 380

Asp Arg Phe Phe Ile Arg Tyr Glu Ala Phe Asp Leu His Gly Thr Ser
385 390 395 400

Leu Gly Ser Asn Ile Phe Pro Lys Pro Ser Val Ile His Asp Glu Gly
405 410 415

Ala

<210> 35

<211> 415

<212> PRT

<213> Haliotis tuberculata

<400> 35

Gly His His Gln Ala Asp Glu Tyr Asp Glu Val Val Thr Ala Ala Ser
1 5 10 15

His Ile Arg Lys Asn Leu Lys Asp Leu Ser Lys Gly Glu Val Glu Ser
20 25 30

Leu Arg Ser Ala Phe Leu Gln Leu Gln Asn Asp Gly Val Tyr Glu Asn
35 40 45

Ile Ala Lys Phe His Gly Lys Pro Gly Leu Cys Asp Asp Asn Gly Arg
50 55 60

Lys Val Ala Cys Cys Val His Gly Met Pro Thr Phe Pro Gln Trp His
65 70 75 80

Arg Leu Tyr Val Leu Gln Val Glu Asn Ala Leu Leu Glu Arg Gly Ser
85 90 95

Ala Val Ser Val Pro Tyr Trp Asp Trp Thr Glu Thr Phe Thr Glu Leu
100 105 110

Pro Ser Leu Ile Ala Glu Ala Thr Tyr Phe Asn Ser Arg Gln Gln Thr
115 120 125

Phe Asp Pro Asn Pro Phe Phe Arg Gly Lys Ile Ser Phe Glu Asn Ala
130 135 140

28

Val Thr Thr Arg Asp Pro Gln Pro Glu Leu Tyr Val Asn Arg Tyr Tyr
 145 150 155 160
 Tyr Gln Asn Val Met Leu Val Phe Glu Gln Asp Asn Tyr Cys Asp Phe
 165 170 175
 Glu Ile Gln Phe Glu Met Val His Asn Val Leu His Ala Trp Leu Gly
 180 185 190
 Gly Arg Ala Thr Tyr Ser Ile Ser Ser Leu Asp Tyr Ser Ala Phe Asp
 195 200 205
 Pro Val Phe Phe Leu His His Ala Asn Thr Asp Arg Leu Trp Ala Ile
 210 215 220
 Trp Gln Glu Leu Gln Arg Tyr Arg Lys Lys Pro Tyr Asn Glu Ala Asp
 225 230 235 240
 Cys Ala Ile Asn Leu Met Arg Lys Pro Leu His Pro Phe Asp Asn Ser
 245 250 255
 Asp Leu Asn His Asp Pro Val Thr Phe Lys Tyr Ser Lys Pro Thr Asp
 260 265 270
 Gly Phe Asp Tyr Gln Asn Asn Phe Gly Tyr Lys Tyr Asp Asn Leu Glu
 275 280 285
 Phe Asn His Phe Ser Ile Pro Arg Leu Glu Glu Ile Ile Arg Ile Arg
 290 295 300
 Gln Arg Gln Asp Arg Val Phe Ala Gly Phe Leu Leu His Asn Ile Gly
 305 310 315 320
 Thr Ser Ala Thr Val Glu Ile Phe Val Cys Val Pro Thr Thr Ser Gly
 325 330 335
 Glu Gln Asn Cys Glu Asn Lys Ala Gly Thr Phe Ala Val Leu Gly Gly
 340 345 350
 Glu Thr Glu Met Ala Phe His Phe Asp Arg Leu Tyr Arg Phe Asp Ile
 355 360 365
 Ser Glu Thr Leu Arg Asp Leu Gly Ile Gln Leu Asp Ser His Asp Phe
 370 375 380
 Asp Leu Ser Ile Lys Ile Gln Gly Val Asn Gly Ser Tyr Leu Asp Pro
 385 390 395 400
 His Ile Leu Pro Glu Pro Ser Leu Ile Phe Val Pro Gly Ser Ser
 405 410 415

<210> 36

<211> 418

<212> PRT

<213> Haliotis tuberculata

29

<400> 36
 Ser Phe Leu Arg Pro Asp Gly His Ser Asp Asp Ile Leu Val Arg Lys
 1 5 10 15
 Glu Val Asn Ser Leu Thr Thr Arg Glu Thr Ala Ser Leu Ile His Ala
 20 25 30
 Leu Lys Ser Met Gln Glu Asp His Ser Pro Asp Gly Phe Gln Ala Ile
 35 40 45
 Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Ser Pro Ser Ala Ala
 50 55 60
 His Arg Tyr Ala Cys Cys Val His Gly Met Ala Thr Phe Pro Gln Trp
 65 70 75 80
 His Arg Leu Tyr Thr Val Gln Phe Gln Asp Ala Leu Arg Arg His Gly
 85 90 95
 Ala Thr Val Gly Val Pro Tyr Trp Asp Trp Leu Arg Pro Gln Ser His
 100 105 110
 Leu Pro Glu Leu Val Thr Met Glu Thr Tyr His Asp Ile Trp Ser Asn
 115 120 125
 Arg Asp Phe Pro Asn Pro Phe Tyr Gln Ala Asn Ile Glu Phe Glu Gly
 130 135 140
 Glu Asn Ile Thr Thr Glu Arg Glu Val Ile Ala Asp Lys Leu Phe Val
 145 150 155 160
 Lys Gly Gly His Val Phe Asp Lys Leu Val Leu Gln Thr Ser His Pro
 165 170 175
 Ser Ala Glu Gln Glu Asn Tyr Cys Asp Phe Glu Ile Gln Phe Glu Ile
 180 185 190
 Leu His Asn Gly Val His Thr Trp Val Gly Gly Ser Arg Thr Tyr Ser
 195 200 205
 Ile Gly His Leu His Tyr Ala Phe Tyr Asp Pro Leu Phe Tyr Leu His
 210 215 220
 His Phe Gln Thr Asp Arg Ile Trp Ala Ile Trp Gln Glu Leu Gln Glu
 225 230 235 240
 Gln Arg Gly Leu Ser Gly Asp Glu Ala His Cys Ala Leu Glu Gln Met
 245 250 255
 Arg Glu Pro Leu Lys Pro Phe Ser Phe Gly Ala Pro Tyr Asn Trp Asn
 260 265 270
 Gln Leu Thr Gln Asp Phe Ser Arg Pro Glu Asp Thr Phe Asp Tyr Arg
 275 280 285
 Lys Phe Gly Tyr Glu Tyr Asp Asn Leu Glu Phe Leu Gly Met Ser Val
 290 295 300

30

Ala Glu Leu Asp Gln Tyr Ile Ile Glu His Gln Glu Asn Asp Arg Val
 305 310 315 320

Phe Ala Gly Phe Leu Leu Ser Gly Phe Gly Gly Ser Ala Ser Val Asn
 325 330 335

Phe Gln Val Cys Arg Ala Asp Ser Thr Cys Gln Asp Ala Gly Tyr Phe
 340 345 350

Thr Val Leu Gly Gly Ser Ala Glu Met Ala Trp Ala Phe Asp Arg Leu
 355 360 365

Tyr Lys Tyr Asp Ile Thr Glu Thr Leu Glu Lys Met His Leu Arg Tyr
 370 375 380

Asp Asp Asp Phe Thr Ile Ser Val Ser Leu Thr Ala Asn Asn Gly Thr
 385 390 395 400

Val Leu Ser Ser Ser Leu Ile Pro Thr Pro Ser Val Ile Phe Gln Arg
 405 410 415

Gly His

<210> 37
 <211> 416
 <212> PRT
 <213> Haliotis tuberculata

<400> 37
 Arg Asp Ile Asn Thr Arg Ser Met Ser Pro Asn Arg Val Arg Arg Glu
 1 5 10 15

Leu Ser Asp Leu Ser Ala Arg Asp Leu Ser Ser Leu Lys Ser Ala Leu
 20 25 30

Arg Asp Leu Gln Glu Asp Asp Gly Pro Asn Gly Tyr Gln Ala Leu Ala
 35 40 45

Ala Phe His Gly Leu Pro Ala Gly Cys His Asp Ser Arg Gly Asn Glu
 50 55 60

Ile Ala Cys Cys Ile His Gly Met Pro Thr Phe Pro Gln Trp His Arg
 65 70 75 80

Leu Tyr Thr Leu Gln Leu Glu Met Ala Leu Arg Arg His Gly Ser Ser
 85 90 95

Val Ala Ile Pro Tyr Trp Asp Trp Thr Lys Pro Ile Ser Glu Leu Pro
 100 105 110

Ser Leu Phe Thr Ser Pro Glu Tyr Tyr Asp Pro Trp His Asp Ala Val
 115 120 125

31

Val Asn Asn Pro Phe Ser Lys Gly Phe Val Lys Phe Ala Asn Thr Tyr
 130 135 140
 Thr Val Arg Asp Pro Gln Glu Met Leu Phe Gln Leu Cys Glu His Gly
 145 150 155 160
 Glu Ser Ile Leu Tyr Glu Gln Thr Leu Leu Ala Leu Glu Gln Thr Asp
 165 170 175
 Tyr Cys Asp Phe Glu Val Gln Phe Glu Val Leu His Asn Val Ile His
 180 185 190
 Tyr Leu Val Gly Gly Arg Gln Thr Tyr Ala Leu Ser Ser Leu His Tyr
 195 200 205
 Ala Ser Tyr Asp Pro Phe Phe Phe Ile His His Ser Phe Val Asp Lys
 210 215 220
 Met Trp Val Val Trp Gln Ala Leu Gln Lys Arg Arg Lys Leu Pro Tyr
 225 230 235 240
 Lys Arg Ala Asp Cys Ala Val Asn Leu Met Thr Lys Pro Met Arg Pro
 245 250 255
 Phe Asp Ser Asp Met Asn Gln Asn Pro Phe Thr Lys Met His Ala Val
 260 265 270
 Pro Asn Thr Leu Tyr Asp Tyr Glu Thr Leu Tyr Tyr Ser Tyr Asp Asn
 275 280 285
 Leu Glu Ile Gly Gly Arg Asn Leu Asp Gln Leu Gln Ala Glu Ile Asp
 290 295 300
 Arg Ser Arg Ser His Asp Arg Val Phe Ala Gly Phe Leu Leu Arg Gly
 305 310 315 320
 Ile Gly Thr Ser Ala Asp Val Arg Phe Trp Ile Cys Arg Asn Glu Asn
 325 330 335
 Asp Cys His Arg Gly Gly Ile Ile Phe Ile Leu Gly Gly Ala Lys Glu
 340 345 350
 Met Pro Trp Ser Phe Asp Arg Asn Phe Lys Phe Asp Ile Thr His Val
 355 360 365
 Leu Glu Asn Ala Gly Ile Ser Pro Glu Asp Val Phe Asp Ala Glu Glu
 370 375 380
 Pro Phe Tyr Ile Lys Val Glu Ile His Ala Val Asn Lys Thr Met Ile
 385 390 395 400
 Pro Ser Ser Val Ile Pro Ala Pro Thr Ile Ile Tyr Ser Pro Gly Glu
 405 410 415

<210> 38
 <211> 402
 <212> PRT
 <213> *Haliotis tuberculata*

<400> 38

Gly Arg Ala Ala Asp Ser Ala His Ser Ala Asn Ile Ala Gly Ser Gly
 1 5 10 15
 Val Arg Lys Asp Val Thr Thr Leu Thr Val Ser Glu Thr Glu Asn Leu
 20 25 30
 Arg Gln Ala Leu Gln Gly Val Ile Asp Asp Thr Gly Pro Asn Gly Tyr
 35 40 45
 Gln Ala Ile Ala Ser Phe His Gly Ser Pro Pro Met Cys Glu Met Asn
 50 55 60
 Gly Arg Lys Val Ala Cys Cys Ala His Gly Met Ala Ser Phe Pro His
 65 70 75 80
 Trp His Arg Leu Tyr Val Lys Gln Met Glu Asp Ala Leu Ala Asp His
 85 90 95
 Gly Ser His Ile Gly Ile Pro Tyr Trp Asp Trp Thr Thr Ala Phe Thr
 100 105 110
 Glu Leu Pro Ala Leu Val Thr Asp Ser Glu Asn Asn Pro Phe His Glu
 115 120 125
 Gly Arg Ile Asp His Leu Gly Val Thr Thr Ser Arg Ser Pro Arg Asp
 130 135 140
 Met Leu Phe Asn Asp Pro Glu Gln Gly Ser Glu Ser Phe Phe Tyr Arg
 145 150 155 160
 Gln Val Leu Leu Ala Leu Glu Gln Thr Asp Tyr Cys Gln Phe Glu Val
 165 170 175
 Gln Phe Glu Leu Thr His Asn Ala Ile His Ser Trp Thr Gly Gly Arg
 180 185 190
 Ser Pro Tyr Gly Met Ser Thr Leu Glu Phe Thr Ala Tyr Asp Pro Leu
 195 200 205
 Phe Trp Leu His His Ser Asn Thr Asp Arg Ile Trp Ala Val Trp Gln
 210 215 220
 Ala Leu Gln Lys Tyr Arg Gly Leu Pro Tyr Asn Glu Ala His Cys Glu
 225 230 235 240
 Ile Gln Val Leu Lys Gln Pro Leu Arg Pro Phe Asn Asp Asp Ile Asn
 245 250 255
 His Asn Pro Ile Thr Lys Thr Asn Ala Arg Pro Ile Asp Ser Phe Asp
 260 265 270

33

Tyr Glu Arg Phe Asn Tyr Gln Tyr Asp Thr Leu Ser Phe His Gly Lys
 275 280 285
 Ser Ile Pro Glu Leu Asn Asp Leu Leu Glu Glu Arg Lys Arg Glu Glu
 290 295 300
 Arg Thr Phe Ala Ala Phe Leu Leu Arg Gly Ile Gly Cys Ser Ala Asp
 305 310 315 320
 Val Val Phe Asp Ile Cys Arg Pro Asn Gly Asp Cys Val Phe Ala Gly
 325 330 335
 Thr Phe Ala Val Leu Gly Gly Glu Leu Glu Met Pro Trp Ser Phe Asp
 340 345 350
 Arg Leu Phe Arg Tyr Asp Ile Thr Arg Val Met Asn Gln Leu His Leu
 355 360 365
 Gln Tyr Asp Ser Asp Phe Ser Phe Arg Val Lys Leu Val Ala Thr Asn
 370 375 380
 Gly Thr Glu Leu Ser Ser Asp Leu Leu Lys Ser Pro Thr Ile Glu His
 385 390 395 400
 Glu Leu

<210> 39

<211> 515

<212> PRT

<213> Haliotis tuberculata

<400> 39

Gly Ala His Arg Gly Pro Val Glu Glu Thr Glu Val Thr Arg Gln His
 1 5 10 15
 Thr Asp Gly Asn Ala His Phe His Arg Lys Glu Val Asp Ser Leu Ser
 20 25 30
 Leu Asp Glu Ala Asn Asn Leu Lys Asn Ala Leu Tyr Lys Leu Gln Asn
 35 40 45
 Asp His Ser Leu Thr Gly Tyr Glu Ala Ile Ser Gly Tyr His Gly Tyr
 50 55 60
 Pro Asn Leu Cys Pro Glu Glu Gly Asp Asp Lys Ile Pro Leu Leu Arg
 65 70 75 80
 Pro Arg Met Gly Ile Phe Pro Tyr Trp His Arg Leu Leu Thr Ile Gln
 85 90 95
 Leu Glu Arg Ala Leu Glu His Asn Gly Ala Leu Leu Gly Val Pro Tyr
 100 105 110

34

Trp	Asp	Trp	Asn	Lys	Asp	Leu	Ser	Ser	Leu	Pro	Ala	Phe	Phe	Ser	Asp
	115						120					125			
Ser	Ser	Asn	Asn	Asn	Pro	Tyr	Phe	Lys	Tyr	His	Ile	Ala	Gly	Val	Gly
	130					135					140				
His	Asp	Thr	Val	Arg	Glu	Pro	Thr	Ser	Leu	Ile	Tyr	Asn	Gln	Pro	Gln
145					150					155					160
Ile	His	Gly	Tyr	Asp	Tyr	Leu	Tyr	Tyr	Leu	Ala	Leu	Thr	Thr	Leu	Glu
				165					170					175	
Glu	Asn	Asn	Tyr	Trp	Asp	Phe	Glu	Val	Gln	Tyr	Glu	Ile	Leu	His	Asn
			180					185					190		
Ala	Val	His	Ser	Trp	Leu	Gly	Gly	Ser	Gln	Lys	Tyr	Ser	Met	Ser	Thr
		195					200					205			
Leu	Glu	Tyr	Ser	Ala	Phe	Asp	Pro	Val	Phe	Met	Ile	Leu	His	Ser	Gly
	210					215					220				
Leu	Asp	Arg	Leu	Trp	Ile	Ile	Trp	Gln	Glu	Leu	Gln	Lys	Ile	Arg	Arg
225					230					235					240
Lys	Pro	Tyr	Asn	Phe	Ala	Lys	Cys	Ala	Tyr	His	Met	Met	Glu	Glu	Pro
				245					250					255	
Leu	Ala	Pro	Phe	Ser	Tyr	Pro	Ser	Ile	Asn	Gln	Asp	Glu	Phe	Thr	Arg
			260					265					270		
Ala	Asn	Ser	Lys	Pro	Ser	Thr	Val	Phe	Asp	Ser	His	Lys	Phe	Gly	Tyr
		275					280					285			
His	Tyr	Asp	Asn	Leu	Asn	Val	Arg	Gly	His	Ser	Ile	Gln	Glu	Leu	Asn
	290					295					300				
Thr	Ile	Ile	Asn	Asp	Leu	Arg	Asn	Thr	Asp	Arg	Ile	Tyr	Ala	Gly	Phe
305					310					315					320
Val	Leu	Ser	Gly	Ile	Gly	Thr	Ser	Ala	Ser	Val	Lys	Ile	Tyr	Leu	Arg
				325					330					335	
Thr	Asp	Asp	Asn	Asp	Glu	Glu	Val	Gly	Thr	Phe	Thr	Val	Leu	Gly	Gly
			340					345					350		
Glu	Arg	Glu	Met	Pro	Trp	Ala	Tyr	Glu	Arg	Val	Phe	Lys	Tyr	Asp	Ile
		355					360					365			
Thr	Glu	Val	Ala	Asp	Arg	Leu	Lys	Ile	Lys	Leu	Trp	Gly	His	Pro	Leu
	370					375					380				
Thr	Ser	Gly	Thr	Gly	Asp	His	Ile	Leu	Thr	Asn	Gly	Ile	Gly	Gly	Lys
385					390					395					400
Gln	Glu	Pro	Thr	Gln	Ile	Leu	Ser	Ser	Ser	Thr	Asp	Leu	Pro	Ile	Met
				405					410					415	

35

Thr Thr Met Phe Leu Leu Ser Gln Xaa Gly Arg Asn Leu His Ile Pro
 420 425 430

Pro Lys Val Val Val Lys Lys Gly Thr Arg Ile Glu Phe His Pro Val
 435 440 445

Asp Asp Ser Val Thr Arg Pro Val Val Asp Leu Gly Ser Tyr Thr Ala
 450 455 460

Leu Phe Asn Cys Val Val Pro Pro Phe Thr Tyr His Gly Phe Glu Leu
 465 470 475 480

Asn His Val Tyr Ser Val Lys Pro Gly Asp Tyr Tyr Val Thr Gly Pro
 485 490 495

Thr Arg Asp Leu Cys Gln Asn Ala Asp Val Arg Ile His Ile His Val
 500 505 510

Glu Asp Glu
 515

<210> 40

<211> 322

<212> PRT

<213> Megathura crenulata

<400> 40

Gly Leu Pro Tyr Trp Asp Trp Thr Glu Pro Met Thr His Ile Pro Gly
 1 5 10 15

Leu Ala Gly Asn Lys Thr Tyr Val Asp Ser His Gly Ala Ser His Thr
 20 25 30

Asn Pro Phe His Ser Ser Val Ile Ala Phe Glu Glu Asn Ala Pro His
 35 40 45

Thr Lys Arg Gln Ile Asp Gln Arg Leu Phe Lys Pro Ala Thr Phe Gly
 50 55 60

His His Thr Asp Leu Phe Asn Gln Ile Leu Tyr Ala Phe Glu Gln Glu
 65 70 75 80

Asp Tyr Cys Asp Phe Glu Val Gln Phe Glu Ile Thr His Asn Thr Ile
 85 90 95

His Ala Trp Thr Gly Gly Ser Glu His Phe Ser Met Ser Ser Leu His
 100 105 110

Tyr Thr Ala Phe Asp Pro Leu Phe Tyr Phe His His Ser Asn Val Asp
 115 120 125

Arg Leu Trp Ala Val Trp Gln Ala Leu Gln Met Arg Arg His Lys Pro
 130 135 140

Tyr Arg Ala His Cys Ala Ile Ser Leu Glu His Met His Leu Lys Pro
 145 150 155 160

36

Phe Ala Phe Ser Ser Pro Leu Asn Asn Asn Glu Lys Thr His Ala Asn
 165 170 175
 Ala Met Pro Asn Lys Ile Tyr Asp Tyr Glu Asn Val Leu His Tyr Thr
 180 185 190
 Tyr Glu Asp Leu Thr Phe Gly Gly Ile Ser Leu Glu Asn Ile Glu Lys
 195 200 205
 Met Ile His Glu Asn Gln Gln Glu Asp Arg Ile Tyr Ala Gly Phe Leu
 210 215 220
 Leu Ala Gly Ile Arg Thr Ser Ala Asn Val Asp Ile Phe Ile Lys Thr
 225 230 235 240
 Thr Asp Ser Val Gln His Lys Ala Gly Thr Phe Ala Val Leu Gly Gly
 245 250 255
 Ser Lys Glu Met Lys Trp Gly Phe Asp Arg Val Phe Lys Phe Asp Ile
 260 265 270
 Thr His Val Leu Lys Asp Leu Asp Leu Thr Ala Asp Gly Asp Phe Glu
 275 280 285
 Val Thr Val Asp Ile Thr Glu Val Asp Gly Thr Lys Leu Ala Ser Ser
 290 295 300
 Leu Ile Pro His Ala Ser Val Ile Arg Glu His Ala Arg Gly Lys Leu
 305 310 315 320
 Asn Arg

<210> 41
 <211> 414
 <212> PRT
 <213> Megathura crenulata

<400> 41
 Val Lys Phe Asp Lys Val Pro Arg Ser Arg Leu Ile Arg Lys Asn Val
 1 5 10 15
 Asp Arg Leu Ser Pro Glu Glu Met Asn Glu Leu Arg Lys Ala Leu Ala
 20 25 30
 Leu Leu Lys Glu Asp Lys Ser Ala Gly Gly Phe Gln Gln Leu Gly Ala
 35 40 45
 Phe His Gly Glu Pro Lys Trp Cys Pro Ser Pro Glu Ala Ser Lys Lys
 50 55 60
 Phe Ala Cys Cys Val His Gly Met Ser Val Phe Pro His Trp His Arg
 65 70 75 80

37

Leu Leu Thr Val Gln Ser Glu Asn Ala Leu Arg Arg His Gly Tyr Asp
85 90 95

Gly Ala Leu Pro Tyr Trp Asp Trp Thr Ser Pro Leu Asn His Leu Pro
100 105 110

Glu Leu Ala Asp His Glu Lys Tyr Val Asp Pro Glu Asp Gly Val Glu
115 120 125

Lys His Asn Pro Trp Phe Asp Gly His Ile Asp Thr Val Asp Lys Thr
130 135 140

Thr Thr Arg Ser Val Gln Asn Lys Leu Phe Glu Gln Pro Glu Phe Gly
145 150 155 160

His Tyr Thr Ser Ile Ala Lys Gln Val Leu Leu Ala Leu Glu Gln Asp
165 170 175

Asn Phe Cys Asp Phe Glu Ile Gln Tyr Glu Ile Ala His Asn Tyr Ile
180 185 190

His Ala Leu Val Gly Gly Ala Gln Pro Tyr Gly Met Ala Ser Leu Arg
195 200 205

Tyr Thr Ala Phe Asp Pro Leu Phe Tyr Leu His His Ser Asn Thr Asp
210 215 220

Arg Ile Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Lys Pro
225 230 235 240

Tyr Asn Val Ala Asn Cys Ala Val Thr Ser Met Arg Glu Pro Leu Gln
245 250 255

Pro Phe Gly Leu Ser Ala Asn Ile Asn Thr Asp His Val Thr Lys Glu
260 265 270

His Ser Val Pro Phe Asn Val Phe Asp Tyr Lys Thr Asn Phe Asn Tyr
275 280 285

Glu Tyr Asp Thr Leu Glu Phe Asn Gly Leu Ser Ile Ser Gln Leu Asn
290 295 300

Lys Lys Leu Glu Ala Ile Lys Ser Gln Asp Arg Phe Phe Ala Gly Phe
305 310 315 320

Leu Leu Ser Gly Phe Lys Lys Ser Ser Leu Val Lys Phe Asn Ile Cys
325 330 335

Thr Asp Ser Ser Asn Cys His Pro Ala Gly Glu Phe Tyr Leu Leu Gly
340 345 350

Asp Glu Asn Glu Met Pro Trp Ala Tyr Asp Arg Val Phe Lys Tyr Asp
355 360 365

Ile Thr Glu Lys Leu His Asp Leu Lys Leu His Ala Glu Asp His Phe
370 375 380

38

Tyr Ile Asp Tyr Glu Val Phe Asp Leu Lys Pro Ala Ser Leu Gly Lys
385 390 395 400

Asp Leu Phe Lys Gln Pro Ser Val Ile His Glu Pro Arg Ile
405 410

<210> 42

<211> 411

<212> PRT

<213> Megathura crenulata

<400> 42

Gly His His Glu Gly Glu Val Tyr Gln Ala Glu Val Thr Ser Ala Asn
1 5 10 15

Arg Ile Arg Lys Asn Ile Glu Asn Leu Ser Leu Gly Glu Leu Glu Ser
20 25 30

Leu Arg Ala Ala Phe Leu Glu Ile Glu Asn Asp Gly Thr Tyr Glu Ser
35 40 45

Ile Ala Lys Phe His Gly Ser Pro Gly Leu Cys Gln Leu Asn Gly Asn
50 55 60

Pro Ile Ser Cys Cys Val His Gly Met Pro Thr Phe Pro His Trp His
65 70 75 80

Arg Leu Tyr Val Val Val Val Glu Asn Ala Leu Leu Lys Lys Gly Ser
85 90 95

Ser Val Ala Val Pro Tyr Trp Asp Trp Thr Lys Arg Ile Glu His Leu
100 105 110

Pro His Leu Ile Ser Asp Ala Thr Tyr Tyr Asn Ser Arg Gln His His
115 120 125

Tyr Glu Thr Asn Pro Phe His His Gly Lys Ile Thr His Glu Asn Glu
130 135 140

Ile Thr Thr Arg Asp Pro Lys Asp Ser Leu Phe His Ser Asp Tyr Phe
145 150 155 160

Tyr Glu Gln Val Leu Tyr Ala Leu Glu Gln Asp Asn Phe Cys Asp Phe
165 170 175

Glu Ile Gln Leu Glu Ile Leu His Asn Ala Leu His Ser Leu Leu Gly
180 185 190

Gly Lys Gly Lys Tyr Ser Met Ser Asn Leu Asp Tyr Ala Ala Phe Asp
195 200 205

Pro Val Phe Phe Leu His His Ala Thr Thr Asp Arg Ile Trp Ala Ile
210 215 220

Trp Gln Asp Leu Gln Arg Phe Arg Lys Arg Pro Tyr Arg Glu Ala Asn
225 230 235 240

39

```
<210> 43
<211> 111
<212> PRT
<213> Megathura crenulata
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<400> 43

Asp Ser Ala His Thr Asp Asp Gly His Thr Glu Pro Val Met Ile Arg
1 5 10 15

Lys Asp Ile Thr Gln Leu Asp Lys Arg Gln Gln Leu Ser Leu Val Lys
20 25 30

Ala Leu Glu Ser Met Lys Ala Asp His Ser Ser Asp Gly Phe Gln Ala
35 40 45

Ile Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Ser Pro Ala Ala
50 55 60

Ser Lys Arg Phe Ala Cys Cys Val His Gly Met Pro Thr Phe Pro Gln
65 70 75 80

40

Trp His Arg Leu Tyr Thr Val Gln Phe Gln Asp Ser Leu Arg Lys His
85 90 95

Gly Ala Val Val Gly Leu Pro Tyr Trp Asp Trp Thr Leu Pro Arg
100 105 110

<210> 44
<211> 317
<212> PRT
<213> Megathura crenulata

<400> 44

Gly Leu Pro Tyr Trp Asp Trp Thr Met Pro Met Ser His Leu Pro Glu
1 5 10 15

Leu Ala Thr Ser Glu Thr Tyr Leu Asp Pro Val Thr Gly Glu Thr Lys
20 25 30

Asn Asn Pro Phe His His Ala Gln Val Ala Phe Glu Asn Gly Val Thr
35 40 45

Ser Arg Asn Pro Asp Ala Lys Leu Phe Met Lys Pro Thr Tyr Gly Asp
50 55 60

His Thr Tyr Leu Phe Asp Ser Met Ile Tyr Ala Phe Glu Gln Glu Asp
65 70 75 80

Phe Cys Asp Phe Glu Val Gln Tyr Glu Leu Thr His Asn Ala Ile His
85 90 95

Ala Trp Val Gly Gly Ser Glu Lys Tyr Ser Met Ser Ser Leu His Tyr
100 105 110

Thr Ala Phe Asp Pro Ile Phe Tyr Leu His His Ser Asn Val Asp Arg
115 120 125

Leu Trp Ala Ile Trp Gln Ala Leu Gln Ile Arg Arg Gly Lys Ser Tyr
130 135 140

Lys Ala His Cys Ala Ser Ser Gln Glu Arg Glu Pro Leu Lys Pro Phe
145 150 155 160

Ala Phe Ser Ser Pro Leu Asn Asn Asn Glu Lys Thr Tyr His Asn Ser
165 170 175

Val Pro Thr Asn Val Tyr Asp Tyr Val Gly Val Leu His Tyr Arg Tyr
180 185 190

Asp Asp Leu Gln Phe Gly Gly Met Thr Met Ser Glu Leu Glu Glu Tyr
195 200 205

Ile His Lys Gln Thr Gln His Asp Arg Thr Phe Ala Gly Phe Phe Leu
210 215 220

Ser Tyr Ile Gly Thr Ser Ala Ser Val Asp Ile Phe Ile Asn Arg Glu
225 230 235 240


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<400> 45
Asp Ala Lys Asp Phe Gly His Ser Arg Lys Ile Arg Lys Ala Val Asp
 1          5          10          15
Ser Leu Thr Val Glu Glu Gln Thr Ser Leu Arg Arg Ala Met Ala Asp
          20          25          30
Leu Gln Asp Asp Lys Thr Ser Gly Gly Phe Gln Gln Ile Ala Ala Phe
          35          40          45
His Gly Glu Pro Lys Trp Cys Pro Ser Pro Glu Ala Glu Lys Lys Phe
          50          55          60
Ala Cys Cys Val His Gly Met Ala Val Phe Pro His Trp His Arg Leu
 65          70          75          80
Leu Thr Val Gln Gly Glu Asn Ala Leu Arg Lys His Gly Phe Thr Gly
          85          90          95
Gly Leu Pro Tyr Trp Asp Trp Thr Arg Ser Met Ser Ala Leu Pro His
          100          105          110
he Val Ala Asp Pro Thr Tyr Asn Asp Ala Ile Ser Ser Gln Glu Glu
          115          120          125
sp Asn Pro Trp His His Gly His Ile Asp Ser Val Gly His Asp Thr
          130          135          140
hr Arg Asp Val Arg Asp Asp Leu Tyr Gln Ser Pro Gly Phe Gly His
 45          150          155          160
yr Thr Asp Ile Ala Gln Gln Val Leu Leu Ala Phe Glu Gln Asp Ser
          165          170          175

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42

Phe Cys Asp Phe Glu Val Gln Phe Glu Ile Ala His Asn Phe Ile His
 180 185 190
 Ala Leu Ile Gly Gly Asn Glu Pro Tyr Ser Met Ser Ser Leu Arg Tyr
 195 200 205
 Thr Thr Tyr Asp Pro Ile Phe Phe Leu His His Ser Ser Thr Asp Arg
 210 215 220
 Leu Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Lys Pro Tyr
 225 230 235 240
 Asn Thr Ala Asn Cys Ala Ile Ala Ser Met Arg Lys Pro Leu Gln Pro
 245 250 255
 Phe Gly Leu Asp Ser Val Ile Asn Pro Asp Asp Glu Thr Arg Glu His
 260 265 270
 Ser Val Pro Phe Arg Val Phe Asp Tyr Lys Asn Asn Phe Asp Tyr Glu
 275 280 285
 Tyr Glu Ser Leu Ala Phe Asn Gly Leu Ser Ile Ala Gln Leu Asp Arg
 290 295 300
 Glu Leu Gln Arg Arg Lys Ser His Asp Arg Val Phe Ala Gly Phe Leu
 305 310 315 320
 Leu His Glu Ile Gly Gln Ser Ala Lys His Asn Val Ser Asp Cys Asp
 325 330 335
 His Tyr Ala Gly Glu Phe Tyr Ile Leu Gly Asp Glu Ala Glu Met Pro
 340 345 350
 Trp Arg Tyr Asp Arg Val Tyr Lys Tyr Glu Ile Thr Gln Gln Leu His
 355 360 365
 Asp Leu Asp Leu His Val Gly Asp Asn Phe Phe Leu Lys Tyr Glu Ala
 370 375 380
 Phe Asp Leu Asn Gly Gly Ser Leu Gly Gly Ser Ile Phe Ser Gln Pro
 385 390 395 400
 Ser Val Ile Phe Glu Pro Ala Ala Gly Met Phe
 405 410

<210> 46

<211> 109

<212> PRT

<213> Megathura crenulata

<400> 46

Gly Ser His Gln Ala Asp Glu Tyr Arg Glu Ala Val Thr Ser Ala Ser
 1 5 10 15
 His Ile Arg Lys Asn Ile Arg Asp Leu Ser Glu Gly Glu Ile Glu Ser
 20 25 30

43

Ile Arg Ser Ala Phe Leu Gln Ile Gln Lys Glu Gly Ile Tyr Glu Asn
 35 40 45

Ile Ala Lys Phe His Gly Lys Pro Gly Leu Cys Glu His Asp Gly His
 50 55 60

Pro Val Ala Cys Cys Val His Gly Met Pro Thr Phe Pro His Trp His
 65 70 75 80

Arg Leu Tyr Val Leu Gln Val Glu Asn Ala Leu Leu Glu Arg Gly Ser
 85 90 95

Ala Val Ala Val Pro Tyr Trp Asp Trp Thr Leu Pro Arg
 100 105

<210> 47

<211> 329

<212> PRT

<213> Megathura crenulata

<400> 47

Met Ala Val Phe Pro His Trp His Arg Leu Phe Val Lys Gln Met Glu
 1 5 10 15

Asp Ala Leu Ala Ala His Gly Ala His Ile Gly Ile Pro Tyr Trp Asp
 20 25 30

Trp Thr Ser Ala Phe Ser His Leu Pro Ala Leu Val Thr Asp His Glu
 35 40 45

Asn Asn Pro Phe His His Gly His Ile Gly His Leu Asn Val Asp Thr
 50 55 60

Ser Arg Ser Pro Arg Asp Met Leu Phe Asn Asp Pro Glu Gln Gly Ser
 65 70 75 80

Glu Ser Phe Phe Tyr Arg Gln Val Leu Leu Thr Leu Glu Gln Thr Asp
 85 90 95

Phe Cys Gln Phe Glu Val Gln Phe Glu Leu Thr His Asn Ala Ile His
 100 105 110

Ser Trp Thr Gly Gly His Thr Pro Tyr Gly Met Ser Ser Leu Glu Tyr
 115 120 125

Thr Ala Tyr Asp Pro Leu Phe Tyr Leu His His Ser Asn Thr Asp Arg
 130 135 140

Ile Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Leu Pro Tyr
 145 150 155 160

Asn Ala Ala His Cys Asp Ile Gln Val Leu Lys Gln Pro Leu Lys Pro
 165 170 175

44

Phe Ser Glu Ser Arg Asn Pro Asn Pro Val Thr Arg Ala Asn Ser Arg
 180 185 190
 Ala Val Asp Ser Phe Asp Tyr Glu Lys Phe Asn Tyr Gln Tyr Asp Thr
 195 200 205
 Leu Thr Phe His Gly Leu Ser Ile Pro Glu Leu Asp Ala Met Leu Gln
 210 215 220
 Glu Arg Lys Lys Glu Glu Arg Thr Phe Ala Ala Phe Leu Leu His Gly
 225 230 235 240
 Phe Gly Ala Ser Ala Asp Val Ser Phe Asp Val Cys Thr Pro Asp Gly
 245 250 255
 His Cys Ala Phe Ala Gly Thr Phe Ala Val Leu Gly Gly Glu Leu Glu
 260 265 270
 Met Pro Trp Ser Phe Glu Arg Leu Phe Arg Tyr Asp Ile Thr Lys Val
 275 280 285
 Leu Lys Gln Met Asn Leu His Tyr Asp Ser Glu Phe His Phe Glu Leu
 290 295 300
 Lys Ile Val Gly Thr Asp Gly Thr Glu Leu Pro Ser Asp Arg Ile Lys
 305 310 315 320
 Ser Pro Thr Ile Glu His His Gly Gly
 325

<210> 48

<211> 103

<212> PRT

<213> Megathura crenulata

<400> 48

Gly His Asp His Ser Glu Arg His Asp Gly Phe Phe Arg Lys Glu Val
 1 5 10 15
 Gly Ser Leu Ser Leu Asp Glu Ala Asn Asp Leu Lys Asn Ala Leu Tyr
 20 25 30
 Lys Leu Gln Asn Asp Gln Gly Pro Asn Gly Tyr Glu Ser Ile Ala Gly
 35 40 45
 Tyr His Gly Tyr Pro Phe Leu Cys Pro Glu His Gly Glu Asp Gln Tyr
 50 55 60
 Ala Cys Cys Val His Gly Met Pro Val Phe Pro His Trp His Arg Leu
 65 70 75 80
 His Thr Ile Gln Phe Glu Arg Ala Leu Lys Glu His Gly Ser His Leu
 85 90 95
 Gly Leu Pro Tyr Trp Asp Trp
 100

<210> 49
 <211> 1269
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 49
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 gaaaggacgt gagtcacctc acggatgacg aggtgcaagc tctccacggc gccctccatg 120
 acgtcactgc atctacaggg cctctgagtt tcgaagacat aacatcttac catgccgcac 180
 cagcgtcgtg tgactacaag ggacggaaga tcgcctgctg tgtccacggg atgccagtt 240
 tccccctctg gcacagggca tatgtcgtcc aagccgagcg ggcaactgtg tccaaacgga 300
 agactgtcgg aatgccttac tgggactgga cgcaaacgct gactcactta ccatctcttg 360
 tgactgaacc catctacatt gacagtaaag gtggaaaggc tcaaaccaac tactggtacc 420
 gcggcgagat agcgttcac aataagaaga ctgcgcgagc tgtagatgat cgcctattcg 480
 agaaggtgga gcctggtcac tacacacatc ttatggagac tgcctcgcac gctctcgaa 540
 aggacgaatt ctgtaaattt gaaatccagt tcgagttggc tcataatgct atccattact 600
 tggttggcgg taaatttgaa tattcaatgt caaacttgga atacacctcc tacgacccca 660
 tcttcttctt ccaccactcc aacgttgacc gcctcttcgc catctggcag cgtcttcagg 720
 aactgcgagg aaagaatccc aatgcaatgg actgtgcaca tgaactcgct caccagcaac 780
 tccaaccctt caacagggac agcaatccag tccagctcac aaaggaccac tcgacacctg 840
 ctgacctctt tgattacaaa caacttgat acagctacga cagcttaaac ctgaatggaa 900
 tgacgccaga acagctgaaa acagaactag acgaacgcca ctccaaagaa cgtgcgtttg 960
 caagcttccg actcagtggt tttgggggtt ctgccaaagt tgtgtgtctat gcatgtgtcc 1020
 ctgatgatga tccacgcagt gatgactact gcgagaaagc aggcgacttc ttcattcttg 1080
 ggggtcaaag cgaaatgccg tggagattct acagacctt cttctatgat gtaactgaag 1140
 cgggtacatca ctttggagtc ccgctaagt gccaactacta tgtgaaaaca gaactcttca 1200
 gcgtgaatgg cacagcactt tcacctgac ttcttctca accaactggt gcctaccgac 1260
 ctgggaaag 1269

<210> 50
 <211> 569
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 50
 ggtcttccgt actgggactg gacgcagcat ctgactcaac tcccagatct ggtgtcagac 60
 cccttgtttg tcgaccggga aggaggaaag gcccatgaca acgcatggta tcgtggaaac 120
 atcaagtttg agaataagaa gactgcaaga gctgttgacg atcgctttt cgagaaggtt 180
 ggaccaggag agaatacccc actctttgaa ggaattctcg atgctcttga acaggatgaa 240
 ttctgcaact tcgagatcca gtttgagttg gctcacaacg ctatccacta cctggttggc 300
 ggccgtcaca cgtactccat gtctcatctc gagttacacc ctctacgac cccctcttct 360
 tcctccatca ctccaacacc ggaccgcatc ttgcgcatct gggaacgtct tcagggtactc 420
 agaggaaagg accccaacac cgccgactgc gcacacaacc tcatccatga gcccatggaa 480
 ccgttccgtc gggactcgaa ccctcttgac ctcaccaggg aaaactccaa accaattgac 540
 agctttgatt atgcccacct tggctacca 569

<210> 51
 <211> 1246
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 51
 gttacagagg cccagctcc ctctcggat gctcacctcg ccgtcaggaa ggatatcaac 60
 catctgacac gcgaggaggt gtacgagctg cgcagagcta tggagagatt ccaggccgac 120
 acatccgttg atgggtacca ggctacggtt gagtatcacg gcttacctgc tcgatgtcca 180

46

ttccccgagg	ccacaaatag	gttcgcctgt	tgcattccacg	gcatggcgac	attccctcat	240
tggcacagac	tggttcgtcac	ccaggtggaa	gatgctctga	tcaggcgagg	atcgctata	300
gggggtcccct	actgggactg	gactcagcct	atggcgcatc	tcccaggact	tgcagacaac	360
gccacctata	gagatcccat	cagcggggac	agcagacaca	accccttcca	cgatgttgaa	420
gttgccctttg	aaaatggacg	tacagaacgt	caccagata	gtagattgtt	tgaacaacct	480
ttatttgga	aacatacgcg	tctcttcgac	agtatagtct	atgcttttga	gcaggaggac	540
ttctgcgatt	ttgaagttca	atcttgatg	accataata	atattcacgc	ctggattggt	600
ggcggcgaga	agtattccat	gtcttctcta	cactacacag	ccttcgacct	tatcttctac	660
cttcgtcact	ccaacactga	cgggtcttgg	gcaatttggc	aagcgttgca	gatacgaaga	720
aacaggcctt	acaaggctca	ttgtgcttgg	tctgaggaac	gccagcctct	caaaccttct	780
gccttcagtt	ccccactgaa	caacaacgaa	aaaacctacg	aaaactcggg	gcccaccaac	840
gtttacgact	acgaaggagt	ccttggttat	acttatgatg	acctcaactt	cgggggcatg	900
gacctgggtc	agcttgagga	atacatccag	aggcagagac	agagagacag	gacctttgct	960
ggtttctttc	tggtcacatat	tggtacatca	gcgaatgttg	aatcatttat	agaccatggg	1020
actcttcata	cctccgtggg	cacgttttgc	gttcttggcg	gagagaagga	gatgaaatgg	1080
ggatttgacc	gtttgtacaa	atatgagatt	acagatgaac	tgaggcaact	taatctccgt	1140
gctgatgatg	ttttcagcat	ctctgttaaa	gtaactgatg	ttgatggcag	tgagctgtcc	1200
tctgaactca	tcccatctgc	tgctatcatc	ttcgaacgaa	gccata		1246

<210> 52

<211> 1242

<212> DNA

<213> *Haliotis tuberculata*

<400> 52

gtcaccatca	ggctgacgag	tacgacgaag	ttgtaactgc	tgcaagccac	atcagaaaga	60
attttaaaga	tctgtcaaag	ggagaagtag	agagcctaag	gtctgccttc	ctgcaacttc	120
agaacgacgg	agtctatgag	aatattgcca	agttccacgg	caagcctggg	ttgtgtgatg	180
ataacggctcg	caaggttgcc	tggtgtgtcc	atggaatgcc	caccttcccc	cagtggcaca	240
ggctctatgt	cctccaggtg	gagaatgctt	tgctggagag	aggatctgcc	gtctctgtgc	300
catactggga	ctggactgaa	acatttacag	agctgccatc	tttgattgct	gaggctacct	360
atttcaattc	cgtcaacaa	acgtttgacc	ctaactctt	cttcagaggt	aaaatcagtt	420
ttgagaatgc	tggtacaaca	cgtgatcccc	agcctgagct	gtacgttaac	aggtaact	480
acaaaaacgt	catgttggtt	tttgaacagg	acaactactg	cgacttcgag	atacagtttg	540
agatggttca	caatgttctc	catgcttggc	ttggtggaag	agctacttat	tctatttctt	600
ctcttgatta	ttctgcattc	gacctgtgtg	ttttccttca	ccatgcgaac	acagatagat	660
tgtgggccat	ctggcaggag	ctgcagaggt	acaggaagaa	gccatacaat	gaagcggatt	720
gtgccattaa	cctaattgcg	aaacctctac	atcccttcga	caacagtgat	ctcaatcatg	780
atcctgtaac	ctttaaatat	tcaaaaacca	ctgatggctt	tgactaccag	aacaactttg	840
gatacaagta	tgacaacctt	gagttcaatc	atttcagtat	tcccaggctt	gaagaaatca	900
ttcgtattag	acaacgtcaa	gatcgtgtgt	ttgcaggatt	cctccttcac	aacattggga	960
catccgcaac	tggttgagata	ttcgtctgtg	tccctaccac	cagcgggtgag	caaaactgtg	1020
aaaacaaagc	cggaaacattt	gccgtactcg	gaggagaaac	agagatggcg	tttcattttg	1080
acagactcta	caggtttgac	atcagtgaac	cactgaggga	cctcggcata	cagctggaca	1140
gccatgactt	tgacctcagc	atcaagattc	aaggagttaa	tggatcctac	cttgatccac	1200
acatcctgcc	agagccatcc	ttgatttttg	tgcttggttc	aa		1242

<210> 53

<211> 1257

<212> DNA

<213> *Haliotis tuberculata*

<400> 53

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gcctgacaac	caggagact	gcattctctga	tccatgctct	gaaaagtatg	caggaagacc	120
attcacctga	cgggttccaa	gccattgctt	ctttccatgc	tctgccacca	ctctgccctt	180
cacctgtacg	agctcaccgt	tatgcttgct	gtgtccacgg	catggctaca	tttccccagt	240

E R S A T Z B L A T T (REGEL 26)

47

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ggcacagatt gtacactgta cagttccagg atgcactgag gagacatgga gctacggtag 300
gtgtaccgta ttgggattgg ctgcgaccgc agtctcacct accagagctt gtcaccatgg 360
agacatacca tgatatttgg agtaacagag atttcccaa tcctttctac caagccaata 420
ttgagtttga aggagaaaac attacaacag agagagaaat cattgcagac aaactttttg 480
tcaaaggtgg acacgttttt gataaactgg ttcttcaaac aagccatcct agcgctgagc 540
aggaaaacta ctgtgacttt gagattcagt ttgaaattct tcacaacggc gttcacacgt 600
gggtcggagg cagtcgtacc tactctatcg gacatcttca ttacgcattc tacgaccctc 660
ttttctacct tcaccatttc cagacagacc gtatttgggc aatctggcaa gaactccagg 720
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tgaagccttt cagcttcggc gctccttata actggaatca gctcacacag gatttctccc 840
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tgacacctcg atatgatgat gacttcacaa tctctgtcag tctgaccgcc aacaacggaa 1200
ctgtcctgag cagcagtcta atcccaacac cgagtgtcat attccagcgg ggacatc 1257

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<210> 54

<211> 1257

<212> DNA

<213> Megathura crenulata

<400> 54

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attctgcccc cacagatgat ggacacactg aaccagtgat gattcgcaa gatatcacac 60
aattggacaa gcgtcaacaa ctgtcactgg tgaaagccct cgagtccatg aaagccgacc 120
attcatctga tgggttccag gcaatcgctt ccttccatgc tcttctctct ctttgtccat 180
caccagctgc ttcaaagagg ttgctgtgct gcgtccatgg catggcaacg tcccacaaat 240
ggcaccgtct gtacacagtc caattccaag attctctcag aaaacatggg gcagtcgttg 300
gacttccgta ctgggactgg accctacctc gttctgaatt accagagctc ctgaccgtct 360
caactattca tgaccgggag acaggcagag atataccaaa tccatttatt ggttctaaaa 420
tagagtttga aggagaaaac gtacatacta aaagagatat caatagggat cgtctcttcc 480
agggatcaac aaaaacacat cataactggg ttattgagca agcactgctt gctcttgaac 540
aaaccaacta ctgcgacttc gaggttcagt ttgaaattat gcataatggg gttcatacct 600
gggttggagg caaggagccc tatggaattg gccatctgca ttatgcttcc tatgatccac 660
ttttctacat ccatcactcc caaactgatc gtatttgggc tatatggcaa tcgttgacgc 720
gtttcagagg actttctgga tctgaggcta actgtgctgt aaatctcatg aaaactcctc 780
tgaagccttt cagcttttga gcaccatata atcttaatga tcacacgcat gatttctcaa 840
agcctgaaga tacattcgac taccaaaagt ttggatacat atatgacact ctggaatttg 900
caggggtgtc aattcgtggc attgaccata ttgtccgtaa caggcaggaa cattcaaggg 960
tctttgccgg attcttgctt gaaggatttg gcacctctgc cactgtcgat ttccaggtct 1020
gtcgcacagc ggggagactgt gaagatgcag ggtacttcac cgtgttggga ggtgaaaaag 1080
aatgccttg ggcctttgat cggctttaca agtacgacat aacagaaacc ttagacaaga 1140
tgaaccttcg acatgacgaa atcttccaga ttgaagtaac cattacatcc tacgatggaa 1200
ctgtactcga tagtggcctt attcccacac cgtcaatcat ctatgatcct gctcatc 1257

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<210> 55

<211> 1254

<212> DNA

<213> Megathura crenulata

<400> 55

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cagatggttt tgctgccatt gcacccctcc atggtctgcc tgccaaatgt aatgacagcc 180
acaataacga ggtggcatgc tgtatccatg gaatgcctac attccccac tggcagagac 240
tctacacctt ccaatttgag caagctctaa gaagacatgg ctctagtga gcagtacct 300

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48

actgggactg	gacaaagcca	atacataata	ttccacatct	gttcacagac	aaagaataact	360
acgatgtctg	gagaaataaaa	gtaatgccaa	atccatttgc	ccgaggggtat	gtccccctcac	420
acgatacata	cacggtaaga	gacgtccaag	aaggcctggt	ccacctgaca	tcaacgggtg	480
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cctttgtaga	caaggtttgg	gctgtctggc	aggctcttca	agaaaagaga	ggccttccat	720
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gaattagaac	ctcagctgat	gtccaattcc	agatttgtaa	aacatcagaa	gattgtcacc	1020
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tattcaagta	cgatattacc	catgctcttc	atgacgcaca	catcactcca	gaagacgtat	1140
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<210> 56

<211> 509

<212> DNA

<213> Megathura crenulata

<400> 56

accatcacga	agatcatcat	tcttcttcta	tggttgagca	tggtgtcaga	aaggaaatca	60
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accacggtcc	aaatggatac	caggctatag	cagcggtcca	tggaacacca	ccaatgtgcc	180
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aagaggacaa	tcctttccat	catggtcaca	tagactatgt	gggagtggat	acaactcggg	420
cgccccgaga	caagttgttc	aatgatccag	agcgaggatc	agaatcgttc	ttctacaggc	480
aggttctctt	ggctttggag	cagacagat				509

<210> 57

<211> 943

<212> DNA

<213> Megathura crenulata

<400> 57

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gtggcggttg	aaaatggtgt	aacaagcagg	aatcctgatg	ccaaactttt	tatgaaacca	180
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ctccatcact	caaatgttga	tcgtctctgg	gccatttggc	aagctcttca	aatcaggaga	420
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gtttatgact	atgtgggagt	tttgacttat	cgatatgatg	accttcagtt	tgccgggtatg	600
accatgtcag	aacttgagga	atatattcac	aagcagacac	aacatgatag	aacttttgca	660
ggattcttcc	ttcatatat	tggaacatca	gcaagcgtag	atatcttcat	caatcgagaa	720
ggtcacgata	aatacaaaagt	gggaagtttt	gtagtacttg	gtggatccaa	agaaatgaaa	780
tggggctttg	atagaatgta	caagtatgag	atcactgagg	ctctgaagac	gctgaatgtt	840
gcagtggatg	atgggttcag	cattactggt	gagatcaccg	atgttgatgg	atctccccca	900
tctgcagatc	tcattccacc	tcctgctata	atctttgaac	gtg		943

<210> 58
 <211> 1248
 <212> DNA
 <213> Megathura crenulata

<400> 58
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 ggggtttcca gcagattgca gcattccacg gagaacaaaa atggtgtcca agccccgaag 180
 cggagaaaaa atttgcatgc tgtgttcatg gaatggctgt tttccctcac tggcacagat 240
 tgctgacagt tcaaggagaa aatgctctga ggaaacatgg ctttactggg ggactgccct 300
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 ccaatacaga ccgactttgg gccatttggc aagctttgca aaaataccgg gggaaccat 720
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 atgttgagaa taatttcttc cttaaatatg aagcctttga tctgaatggc ggaagtcttg 1200
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<210> 59
 <211> 1257
 <212> DNA
 <213> Megathura crenulata

<400> 59
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 aaaaagaggg tatatatgaa aacattgcaa agttccatgg aaaaccagga ctttgtgaac 180
 atgatggaca tcctgttgct tgttgtgtcc atggcatgcc cacttttccc cactggcaca 240
 gactgtacgt tcttcaggtg gagaatgcgc tcttagaacg aggggtctgca gttgtgttc 300
 cttactggga ctggaccgag aaagctgact ctctgccatc attaatcaat gatgcaactt 360
 atttcaattc acgatcccag acctttgatc ctaatccttt cttcagggga catattgcct 420
 tcgagaatgc tgtgacgtcc agagatcctc agccagaact atgggacaat aaggacttct 480
 acgagaatgt catgctggct cttgagcaag acaacttctg tgactttgag attcagcttg 540
 agctgataca caacgccctt cattctagac ttggaggaag ggctaaatac tccctttcgt 600
 ctcttgatta taccgattt gatcctgtat tttccttca ccatgcaaac gttgacagaa 660
 tctgggccat ctggcaggac ttgcagagat atagaaaagaa accatacaat gaggctgact 720
 gcgcagtcaa cgagatgcgt aaacctcttc aaccatttaa taaccagaa ctttaacagt 780
 attccatgac gcttaaacac aacctccac aagacagttt tgattatcaa aaccgcttca 840
 ggtaccaata tgataacctt caatttaacc acttcagcat acaaaaagcta gaccaaacta 900
 ttcaggctag aaaacaacac gacagagttt ttgctggcct tattcttcac aacattggga 960
 catctgctgt ttagatatt tatatttgcg ttgaacaagg aggagaacaa aactgcaaga 1020
 caaaggcggg ttccttcacg attctggggg gagaaacaga aatgccattc cactttgacc 1080
 gcttgataca atttgacata acgtctgctc tgcataaact tgggtgtccc ttggacggac 1140
 atggattcga catcaaagtt gacgtcagag ctgtcaatgg atcgcatctt gatcaacaca 1200
 tcctcaacga accgagtctg ctttttgttc ctggtgaacg taagaatata tattatg 1257

<210> 60
 <211> 1239
 <212> DNA
 <213> Megathura crenulata

<400> 60
 atgggctttc acaacataat cttgtgcgaa aagaagtaag ctctcttaca acactggaga 60
 aacatttttt gaggaagct ctcaagaaca tgcaagcaga tgattctcca gacggatatc 120
 aagctattgc ttctttccac gctttgcctc ctctttgtcc aagtccatct gctgcacata 180
 gacacgcttg ttgcctccat ggtatggcta ccttccctca gtggcacaga ctctacacag 240
 ttcagttcga agattctttg aaacgacatg gttctattgt cggacttcca tattgggatt 300
 ggctgaaacc gcagtctgca ctccctgatt tgggtgacaca ggagacatac gagcacctgt 360
 tttcacacaa aaccttccca aatccgttcc tcaaggcaaa tatagaattt gagggagagg 420
 gagnaacaac agagagggat gttgatgctg aacacctctt tgcaaaaagga aatctgggtt 480
 acaacaactg gttttgcaat caggcactat atgcactaga acaagaaaat tactgtgact 540
 ttgaaataca gttcgaaatt ttgcataatg gaattcattc atgggttgga ggatcaaaga 600
 cccattcaat aggtcatctt cattacgcat catacgatcc actgttctat atccaccatt 660
 cgagacaga tcgcatttgg gctatctggc aagctctcca ggagcacaga ggtctttcag 720
 ggaaggaagc aactgcgcc ctggagcaaa tgaaagacc tctcaaacct ttcagctttg 780
 gaagtcctta taatttgaac aaacgcactc aagagttctc caagcctgaa gacacatttg 840
 attatcaccg attcgggtat gagtatgatt cctcgaatt tgttggcatg tctgtttcaa 900
 gtttacataa ctatataaaa caacaacagg aagctgatag agtcttcgca ggattccttc 960
 ttaaaggatt tggacaatca gcatccgtat cgtttgatat ctgcagacca gaccagagtt 1020
 gccaagaagc tggatacttc tcagttctcg gtggaagtcc agaaatgccg tggcagtttg 1080
 acaggcttta caagtacgac attacaaaaa cgttgaaaga catgaaactg cgatacgatg 1140
 acacatttac catcaagggt cacataaagg atatagctgg agctgagttg gacagcgatc 1200
 tgattccaac tccttctgtt ctcttgaag aaggaaagc 1239

<210> 61
 <211> 1251
 <212> DNA
 <213> Haliotis tuberculata

<400> 61
 atgggatcaa tgtacgtcac gttggctgta atcggattcg tatggaacta tctgaactca 60
 ccgagagaga tctcgccagc ctgaaatctg caatgaggtc tctacaagct gacgatgggg 120
 tgaacgggta tcaagccatt gcatcattcc acggtctccc ggcttcttgt catgatgatg 180
 agggacatga gattgcctgt tgtatccacg gaatgccagt attcccacac tggcacaggc 240
 tttacacct gcaaattggac atggctctgt tatctcacgg atctgctgtt gctattccat 300
 actgggactg gaccaaacct atcagcaaac tgcctgatct cttaccagc cctgaatatt 360
 acgatccttg gagggatgca gttgtcaata atccatttgc taaaggctac attaaatccg 420
 aggacgctta cacggtttag gatcctcagg acattttgta ccacttgcag gacgaaacgg 480
 gaacatctgt tttgttagat caaactcttt tagccttaga gcagacagat ttctgtgatt 540
 ttgaggttca atttgaggtc gtccataatg ctattcacta cttgggtgggt ggtcgacaag 600
 tttatgctct ttcttctcaa cactatgctt catatgaccc agccttcttt attcatcact 660
 cctttgttga caaaatatgg gcagtctggc aagctctgca aaagaagaga aagcgtccct 720
 atcataaagc ggattgtgct cttaacatga tgaccaaacc aatgcgacca tttgcacacg 780
 atttcaatca caatggattc acaaaaatgc acgcagtcct caacactcta tttgactttc 840
 aggaccttt ctacacgtat gacaacttag aaattgctgg catgaatgtt aatcagttgg 900
 aagcggaaat caaccggcga aaaagccaaa caagagtctt tgccgggttc cttctacatg 960
 gcatttggaag atcagctgat gtacgatttt ggatttgcaa gacagctgac gactgccacg 1020
 catctggcat gatctttatc ttaggaggtt cttaaagagat gcactgggcc tatgacagga 1080
 actttaata cgacatcacc caagctttga aggtcagtc cataccctt gaagatgtgt 1140
 ttgacactga tgctcctttc ttcattaaag tggaggtcca tgggtgtaaac aagactgctc 1200
 tcccatcttc agctatccca gcacctacta taatctactc agctggtgaa g 1251

<210> 62
 <211> 1185
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 62
 atcatattgc tggcagtgga gtcaggaaag acgtgacgtc tcttacgca tctgagatag 60
 agaacctgag gcatgctctg caaagcgtga tggatgatga tggacccaat ggattccagg 120
 caattgctgc ttatcacgga agtcctccca tgtgtcacat gcntgatggt agagacgttg 180
 catgttgtag tcatggaatg gcatctttcc ctactggca cagactgttt gtgaaacaga 240
 tggaggatgc actggctgcg catggagctc acattggcat accatactgg gattggacaa 300
 gtgcgtttag tcatctgcct gccctagtga ctgaccacga gcacaatccc ttccaccacg 360
 gacatattgc tcatcggaat gtggatacat ctcgatctcc gagagacatg ctgttcaatg 420
 accccgaaca cgggtcagaa tcattcttct atagacaggt tctcttggt ctagaacaga 480
 cagacttctg ccaatttgaa gttcagtttg aaataacaca caatgcaatc cactcttgga 540
 ctggaggaca tactccatat ggaatgtcat cactggaata tacagcatat gatccactct 600
 tttatctcca ccattccaac actgatcgta tctgggcat ctggcaggca ctccagaaat 660
 acagaggttt tcaatacaac gcagctcatt gcgatatcca ggttctgaaa caacctctta 720
 aaccattcag cgagtcagg aatccaaacc cagtcaccag agccaattct agggcagtcg 780
 attcatttga ttatgagaga ctcaattatc aatatgacac acttaccttc cacggacatt 840
 ctatctcaga acttgatgcc atgcttcaag agagaaagaa ggaagagaga acatttgcag 900
 ccttctgtt gcacggattt ggcgccagtg ctgatgttcc gtttgatgtc tgcacacctg 960
 atggtcattg tgctttgct ggaaccttcg cggtaacttg tggggagctt gagatgccct 1020
 ggtcctttga aagattgttc cgttacgata tcacaaaggt tctcaagcag atgaatcttc 1080
 actatgattc tgagttccac tttgagttga agattgttgg cacagatgga acagaactgc 1140
 catcggatcg tatcaagagc cctaccattg aacaccatgg aggag 1185

<210> 63
 <211> 422
 <212> PRT
 <213> *Haliotis tuberculata*

<220>
 <221> SIGNAL
 <222> (1)..(15)

<400> 63
 Leu Val Gln Phe Leu Leu Val Ala Leu Val Val Gly Ala Gly Ala Asp
 1 5 10 15
 Asn Val Val Arg Lys Asp Val Ser His Leu Thr Asp Asp Glu Val Gln
 20 25 30
 Ala Leu His Gly Ala Leu His Asp Val Thr Ala Ser Thr Gly Pro Leu
 35 40 45
 Ser Phe Glu Asp Ile Thr Ser Tyr His Ala Ala Pro Ala Ser Cys Asp
 50 55 60
 Tyr Lys Gly Arg Lys Ile Ala Cys Cys Val His Gly Met Pro Ser Phe
 65 70 75 80
 Pro Phe Trp His Arg Ala Tyr Val Val Gln Ala Glu Arg Ala Leu Leu
 85 90 95
 Ser Lys Arg Lys Thr Val Gly Met Pro Tyr Trp Asp Trp Thr Gln Thr
 100 105 110

Leu Thr His Leu Pro Ser Leu Val Thr Glu Pro Ile Tyr Ile Asp Ser
 115 120 125
 Lys Gly Gly Lys Ala Gln Thr Asn Tyr Trp Tyr Arg Gly Glu Ile Ala
 130 135 140
 Phe Ile Asn Lys Lys Thr Ala Arg Ala Val Asp Asp Arg Leu Phe Glu
 145 150 155 160
 Lys Val Glu Pro Gly His Tyr Thr His Leu Met Glu Thr Val Leu Asp
 165 170 175
 Ala Leu Glu Gln Asp Glu Phe Cys Lys Phe Glu Ile Gln Phe Glu Leu
 180 185 190
 Ala His Asn Ala Ile His Tyr Leu Val Gly Gly Lys Phe Glu Tyr Ser
 195 200 205
 Met Ser Asn Leu Glu Tyr Thr Ser Tyr Asp Pro Ile Phe Phe Leu His
 210 215 220
 His Ser Asn Val Asp Arg Leu Phe Ala Ile Trp Gln Arg Leu Gln Glu
 225 230 235 240
 Leu Arg Gly Lys Asn Pro Asn Ala Met Asp Cys Ala His Glu Leu Ala
 245 250 255
 His Gln Gln Leu Gln Pro Phe Asn Arg Asp Ser Asn Pro Val Gln Leu
 260 265 270
 Thr Lys Asp His Ser Thr Pro Ala Asp Leu Phe Asp Tyr Lys Gln Leu
 275 280 285
 Gly Tyr Ser Tyr Asp Ser Leu Asn Leu Asn Gly Met Thr Pro Glu Gln
 290 295 300
 Leu Lys Thr Glu Leu Asp Glu Arg His Ser Lys Glu Arg Ala Phe Ala
 305 310 315 320
 Ser Phe Arg Leu Ser Gly Phe Gly Gly Ser Ala Asn Val Val Val Tyr
 325 330 335
 Ala Cys Val Pro Asp Asp Asp Pro Arg Ser Asp Asp Tyr Cys Glu Lys
 340 345 350
 Ala Gly Asp Phe Phe Ile Leu Gly Gly Gln Ser Glu Met Pro Trp Arg
 355 360 365
 Phe Tyr Arg Pro Phe Phe Tyr Asp Val Thr Glu Ala Val His His Leu
 370 375 380
 Gly Val Pro Leu Ser Gly His Tyr Tyr Val Lys Thr Glu Leu Phe Ser
 385 390 395 400
 Val Asn Gly Thr Ala Leu Ser Pro Asp Leu Leu Pro Gln Pro Thr Val
 405 410 415

Ala Tyr Arg Pro Gly Lys
420

<210> 64

<211> 511

<212> PRT

<213> *Haliotis tuberculata*

<400> 64

Val His Arg Gly Gly Asn His Glu Asp Glu His His Asp Asp Arg Leu
1 5 10 15

Ala Asp Val Leu Ile Arg Lys Glu Val Asp Phe Leu Ser Leu Gln Glu
20 25 30

Ala Asn Ala Ile Lys Asp Ala Leu Tyr Lys Leu Gln Asn Asp Asp Ser
35 40 45

Lys Gly Gly Phe Glu Ala Ile Ala Gly Tyr His Gly Tyr Pro Asn Met
50 55 60

Cys Pro Glu Arg Gly Thr Asp Lys Tyr Pro Cys Cys Val His Gly Met
65 70 75 80

Pro Val Phe Pro His Trp His Arg Leu His Thr Ile Gln Met Glu Arg
85 90 95

Ala Leu Lys Asn His Gly Ser Pro Met Gly Ile Pro Tyr Trp Asp Trp
100 105 110

Thr Lys Lys Met Ser Ser Leu Pro Ser Phe Phe Gly Asp Ser Ser Asn
115 120 125

Asn Asn Pro Phe Tyr Lys Tyr Tyr Ile Arg Gly Val Gln His Glu Thr
130 135 140

Thr Arg Asp Val Asn Gln Arg Leu Phe Asn Gln Thr Lys Phe Gly Glu
145 150 155 160

Phe Asp Tyr Leu Tyr Tyr Leu Thr Leu Gln Val Leu Glu Glu Asn Ser
165 170 175

Tyr Cys Asp Phe Glu Val Gln Tyr Glu Ile Leu His Asn Ala Val His
180 185 190

Ser Trp Leu Gly Gly Thr Gly Gln Tyr Ser Met Ser Thr Leu Glu His
195 200 205

Ser Ala Phe Asp Pro Val Phe Met Ile His His Ser Ser Leu Asp Arg
210 215 220

Ile Trp Ile Leu Trp Gln Lys Leu Gln Lys Ile Arg Met Lys Pro Tyr
225 230 235 240

Tyr Ala Leu Asp Cys Ala Gly Asp Arg Leu Met Lys Asp Pro Leu His
 245 250 255
 Pro Phe Asn Tyr Glu Thr Val Asn Glu Asp Glu Phe Thr Arg Ile Asn
 260 265 270
 Ser Phe Pro Ser Ile Leu Phe Asp His Tyr Arg Phe Asn Tyr Glu Tyr
 275 280 285
 Asp Asn Met Arg Ile Arg Gly Gln Asp Ile His Glu Leu Glu Glu Val
 290 295 300
 Ile Gln Glu Leu Arg Asn Lys Asp Arg Ile Phe Ala Gly Phe Val Leu
 305 310 315 320
 Ser Gly Leu Arg Ile Ser Ala Thr Val Lys Val Phe Ile His Ser Lys
 325 330 335
 Asn Asp Thr Ser His Glu Glu Tyr Ala Gly Glu Phe Ala Val Leu Gly
 340 345 350
 Gly Glu Lys Glu Met Pro Trp Ala Tyr Glu Arg Met Leu Lys Leu Asp
 355 360 365
 Ile Ser Asp Ala Val His Lys Leu His Val Lys Asp Glu Asp Ile Arg
 370 375 380
 Phe Arg Val Val Val Thr Ala Tyr Asn Gly Asp Val Val Thr Thr Arg
 385 390 395 400
 Leu Ser Gln Pro Phe Ile Val His Arg Pro Ala His Val Ala His Asp
 405 410 415
 Ile Leu Val Ile Pro Val Gly Ala Gly His Asp Leu Pro Pro Lys Val
 420 425 430
 Val Val Lys Ser Gly Thr Lys Val Glu Phe Thr Pro Ile Asp Ser Ser
 435 440 445
 Val Asn Lys Ala Met Val Glu Leu Gly Ser Tyr Thr Ala Met Ala Lys
 450 455 460
 Cys Ile Val Pro Pro Phe Ser Tyr His Gly Phe Glu Leu Asp Lys Val
 465 470 475 480
 Tyr Ser Val Asp His Gly Asp Tyr Tyr Ile Ala Ala Gly Thr His Ala
 485 490 495
 Leu Cys Glu Gln Asn Leu Arg Leu His Ile His Val Glu His Glu
 500 505 510

<210> 65

<211> 197

<212> PRT

<213> Haliotis tuberculata

<400> 65
 Gly Leu Pro Tyr Trp Asp Trp Thr Gln His Leu Thr Gln Leu Pro Asp
 1 5 10 15
 Leu Val Ser Asp Pro Leu Phe Val Asp Pro Glu Gly Gly Lys Ala His
 20 25 30
 Asp Asn Ala Trp Tyr Arg Gly Asn Ile Lys Phe Glu Asn Lys Lys Thr
 35 40 45
 Ala Arg Ala Val Asp Asp Arg Leu Phe Glu Lys Val Gly Pro Gly Glu
 50 55 60
 Asn Thr Arg Leu Phe Glu Gly Ile Leu Asp Ala Leu Glu Gln Asp Glu
 65 70 75 80
 Phe Cys Asn Phe Glu Ile Gln Phe Glu Leu Ala His Asn Ala Ile His
 85 90 95
 Tyr Leu Val Gly Gly Arg His Thr Tyr Ser Met Ser His Leu Glu Tyr
 100 105 110
 Thr Ser Tyr Asp Pro Leu Phe Phe Leu His His Ser Asn Pro Asp Arg
 115 120 125
 Ile Phe Ala Ile Trp Glu Arg Leu Gln Val Leu Arg Gly Lys Asp Pro
 130 135 140
 Asn Thr Ala Asp Cys Ala His Asn Leu Ile His Glu Pro Met Glu Pro
 145 150 155 160
 Phe Arg Arg His Glu Pro Met Glu Pro Phe Arg Arg Asp Ser Asn Pro
 165 170 175
 Leu Asp Leu Thr Arg Glu Asn Ser Lys Pro Ile Asp Ser Phe Asp Tyr
 180 185 190
 Ala His Leu Gly Tyr
 195

<210> 66
 <211> 415
 <212> PRT
 <213> *Haliotis tuberculata*

<400> 66
 Val Thr Glu Ala Pro Ala Pro Ser Ser Asp Ala His Leu Ala Val Arg
 1 5 10 15
 Lys Asp Ile Asn His Leu Thr Arg Glu Glu Val Tyr Glu Leu Arg Arg
 20 25 30
 Ala Met Glu Arg Phe Gln Ala Asp Thr Ser Val Asp Gly Tyr Gln Ala
 35 40 45

56

Thr Val Glu Tyr His Gly Leu Pro Ala Arg Cys Pro Phe Pro Glu Ala
 50 55 60

Thr Asn Arg Phe Ala Cys Cys Ile His Gly Met Ala Thr Phe Pro His
 65 70 75 80

Trp His Arg Leu Phe Val Thr Gln Val Glu Asp Ala Leu Ile Arg Arg
 85 90 95

Gly Ser Pro Ile Gly Val Pro Tyr Trp Asp Trp Thr Gln Pro Met Ala
 100 105 110

His Leu Pro Gly Leu Ala Asp Asn Ala Thr Tyr Arg Asp Pro Ile Ser
 115 120 125

Gly Asp Ser Arg His Asn Pro Phe His Asp Val Glu Val Ala Phe Glu
 130 135 140

Asn Gly Arg Thr Glu Arg His Pro Asp Ser Arg Leu Phe Glu Gln Pro
 145 150 155 160

Leu Phe Gly Lys His Thr Arg Leu Phe Asp Ser Ile Val Tyr Ala Phe
 165 170 175

Glu Gln Glu Asp Phe Cys Asp Phe Glu Val Gln Phe Glu Met Thr His
 180 185 190

Asn Asn Ile His Ala Trp Ile Gly Gly Gly Glu Lys Tyr Ser Met Ser
 195 200 205

Ser Leu His Tyr Thr Ala Phe Asp Pro Ile Phe Tyr Leu Arg His Ser
 210 215 220

Asn Thr Asp Arg Leu Trp Ala Ile Trp Gln Ala Leu Gln Ile Arg Arg
 225 230 235 240

Asn Arg Pro Tyr Lys Ala His Cys Ala Trp Ser Glu Glu Arg Gln Pro
 245 250 255

Leu Lys Pro Phe Ala Phe Ser Ser Pro Leu Asn Asn Asn Glu Lys Thr
 260 265 270

Tyr Glu Asn Ser Val Pro Thr Asn Val Tyr Asp Tyr Glu Gly Val Leu
 275 280 285

Gly Tyr Thr Tyr Asp Asp Leu Asn Phe Gly Gly Met Asp Leu Gly Gln
 290 295 300

Leu Glu Glu Tyr Ile Gln Arg Gln Arg Gln Arg Asp Arg Thr Phe Ala
 305 310 315 320

Gly Phe Phe Leu Ser His Ile Gly Thr Ser Ala Asn Val Glu Ile Ile
 325 330 335

Ile Asp His Gly Thr Leu His Thr Ser Val Gly Thr Phe Ala Val Leu
 340 345 350

Gly Gly Glu Lys Glu Met Lys Trp Gly Phe Asp Arg Leu Tyr Lys Tyr
 355 360 365
 57
 Glu Ile Thr Asp Glu Leu Arg Gln Leu Asn Leu Arg Ala Asp Asp Val
 370 375 380
 Phe Ser Ile Ser Val Lys Val Thr Asp Val Asp Gly Ser Glu Leu Ser
 385 390 395 400
 Ser Glu Leu Ile Pro Ser Ala Ala Ile Ile Phe Glu Arg Ser His
 405 410 415
 <210> 67
 <211> 414
 <212> PRT
 <213> *Haliotis tuberculata*
 <400> 67
 Gly His His Gln Ala Asp Glu Tyr Asp Glu Val Val Thr Ala Ala Ser
 1 5 10 15
 His Ile Arg Lys Asn Leu Lys Asp Leu Ser Lys Gly Glu Val Glu Ser
 20 25 30
 Leu Arg Ser Ala Phe Leu Gln Leu Gln Asn Asp Gly Val Tyr Glu Asn
 35 40 45
 Ile Ala Lys Phe His Gly Lys Pro Gly Leu Cys Asp Asp Asn Gly Arg
 50 55 60
 Lys Val Ala Cys Cys Val His Gly Met Pro Thr Phe Pro Gln Trp His
 65 70 75 80
 Arg Leu Tyr Val Leu Gln Val Glu Asn Ala Leu Leu Glu Arg Gly Ser
 85 90 95
 Ala Val Ser Val Pro Tyr Trp Asp Trp Thr Glu Thr Phe Thr Glu Leu
 100 105 110
 Pro Ser Leu Ile Ala Glu Ala Thr Tyr Phe Asn Ser Arg Gln Gln Thr
 115 120 125
 Phe Asp Pro Asn Pro Phe Phe Arg Gly Lys Ile Ser Phe Glu Asn Ala
 130 135 140
 Val Thr Thr Arg Asp Pro Gln Pro Glu Leu Tyr Val Asn Arg Tyr Tyr
 145 150 155 160
 Tyr Gln Asn Val Met Leu Val Phe Glu Gln Asp Asn Tyr Cys Asp Phe
 165 170 175
 Glu Ile Gln Phe Glu Met Val His Asn Val Leu His Ala Trp Leu Gly
 180 185 190
 Gly Arg Ala Thr Tyr Ser Ile Ser Ser Leu Asp Tyr Ser Ala Phe Asp
 195 200 205

Pro Val Phe Phe Leu His His Ala Asn Thr Asp Arg Leu Trp Ala Ile
 210 215 220
 Trp Gln Glu Leu Gln Arg Tyr Arg Lys Lys Pro Tyr Asn Glu Ala Asp
 225 230 235 240
 Cys Ala Ile Asn Leu Met Arg Lys Pro Leu His Pro Phe Asp Asn Ser
 245 250 255
 Asp Leu Asn His Asp Pro Val Thr Phe Lys Tyr Ser Lys Pro Thr Asp
 260 265 270
 Gly Phe Asp Tyr Gln Asn Asn Phe Gly Tyr Lys Tyr Asp Asn Leu Glu
 275 280 285
 Phe Asn His Phe Ser Ile Pro Arg Leu Glu Glu Ile Ile Arg Ile Arg
 290 295 300
 Gln Arg Gln Asp Arg Val Phe Ala Gly Phe Leu Leu His Asn Ile Gly
 305 310 315 320
 Thr Ser Ala Thr Val Glu Ile Phe Val Cys Val Pro Thr Thr Ser Gly
 325 330 335
 Glu Gln Asn Cys Glu Asn Lys Ala Gly Thr Phe Ala Val Leu Gly Gly
 340 345 350
 Glu Thr Glu Met Ala Phe His Phe Asp Arg Leu Tyr Arg Phe Asp Ile
 355 360 365
 Ser Glu Thr Leu Arg Asp Leu Gly Ile Gln Leu Asp Ser His Asp Phe
 370 375 380
 Asp Leu Ser Ile Lys Ile Gln Gly Val Asn Gly Ser Tyr Leu Asp Pro
 385 390 395 400
 His Ile Leu Pro Glu Pro Ser Leu Ile Phe Val Pro Gly Ser
 405 410

<210> 68

<211> 419

<212> PRT

<213> Haliotis tuberculata

<400> 68

Ser Ser Phe Leu Arg Pro Asp Gly His Ser Asp Asp Ile Leu Val Arg
 1 5 10 15

Lys Glu Val Asn Ser Leu Thr Thr Arg Glu Thr Ala Ser Leu Ile His
 20 25 30

Ala Leu Lys Ser Met Gln Glu Asp His Ser Pro Asp Gly Phe Gln Ala
 35 40 45

59

Ile Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Ser Pro Ser Ala
 50 55 60
 Ala His Arg Tyr Ala Cys Cys Val His Gly Met Ala Thr Phe Pro Gln
 65 70 75 80
 Trp His Arg Leu Tyr Thr Val Gln Phe Gln Asp Ala Leu Arg Arg His
 85 90 95
 Gly Ala Thr Val Gly Val Pro Tyr Trp Asp Trp Leu Arg Pro Gln Ser
 100 105 110
 His Leu Pro Glu Leu Val Thr Met Glu Thr Tyr His Asp Ile Trp Ser
 115 120 125
 Asn Arg Asp Phe Pro Asn Pro Phe Tyr Gln Ala Asn Ile Glu Phe Glu
 130 135 140
 Gly Glu Asn Ile Thr Thr Glu Arg Glu Val Ile Ala Asp Lys Leu Phe
 145 150 155 160
 Val Lys Gly Gly His Val Phe Asp Lys Leu Val Leu Gln Thr Ser His
 165 170 175
 Pro Ser Ala Glu Gln Glu Asn Tyr Cys Asp Phe Glu Ile Gln Phe Glu
 180 185 190
 Ile Leu His Asn Gly Val His Thr Trp Val Gly Gly Ser Arg Thr Tyr
 195 200 205
 Ser Ile Gly His Leu His Tyr Ala Phe Tyr Asp Pro Leu Phe Tyr Leu
 210 215 220
 His His Phe Gln Thr Asp Arg Ile Trp Ala Ile Trp Gln Glu Leu Gln
 225 230 235 240
 Glu Gln Arg Gly Leu Ser Gly Asp Glu Ala His Cys Ala Leu Glu Gln
 245 250 255
 Met Arg Glu Pro Leu Lys Pro Phe Ser Phe Gly Ala Pro Tyr Asn Trp
 260 265 270
 Asn Gln Leu Thr Gln Asp Phe Ser Arg Pro Glu Asp Thr Phe Asp Tyr
 275 280 285
 Arg Lys Phe Gly Tyr Glu Tyr Asp Asn Leu Glu Phe Leu Gly Met Ser
 290 295 300
 Val Ala Glu Leu Asp Gln Tyr Ile Ile Glu His Gln Glu Asn Asp Arg
 305 310 315 320
 Val Phe Ala Gly Phe Leu Leu Ser Gly Phe Gly Gly Ser Ala Ser Val
 325 330 335
 Asn Phe Gln Val Cys Arg Ala Asp Ser Thr Cys Gln Asp Ala Gly Tyr
 340 345 350

Phe Thr Val Leu Gly Gly Ser Ala Glu Met Ala Trp Ala Phe Asp Arg
 355 360 365
 Leu Tyr Lys Tyr Asp Ile Thr Glu Thr Leu Glu Lys Met His Leu Arg
 370 375 380
 Tyr Asp Asp Asp Phe Thr Ile Ser Val Ser Leu Thr Ala Asn Asn Gly
 385 390 395 400
 Thr Val Leu Ser Ser Ser Leu Ile Pro Thr Pro Ser Val Ile Phe Gln
 405 410 415
 Arg Gly His

<210> 69
 <211> 378
 <212> PRT
 <213> Megathura crenulata

<400> 69
 Arg Tyr Gln Ala Thr Ala Glu Tyr His Gly Leu Pro Ala Arg Cys Pro
 1 5 10 15
 Arg Pro Asp Ala Lys Asp Arg Tyr Ala Cys Cys Val His Gly Met Pro
 20 25 30
 Ile Phe Pro His Trp His Arg Leu Phe Val Thr Gln Val Glu Asp Ala
 35 40 45
 Leu Val Gly Arg Gly Ala Thr Ile Gly Ile Pro Tyr Trp Asp Trp Thr
 50 55 60
 Glu Pro Met Thr His Ile Pro Gly Leu Ala Gly Asn Lys Thr Tyr Val
 65 70 75 80
 Asp Ser His Gly Ala Ser His Thr Asn Pro Phe His Ser Ser Val Ile
 85 90 95
 Ala Phe Glu Glu Asn Ala Pro His Thr Lys Arg Gln Ile Asp Gln Arg
 100 105 110
 Leu Phe Lys Pro Ala Thr Phe Gly His His Thr Asp Leu Phe Asn Gln
 115 120 125
 Ile Leu Tyr Ala Phe Glu Gln Glu Asp Tyr Cys Asp Phe Glu Val Gln
 130 135 140
 Phe Glu Ile Thr His Asn Thr Ile His Ala Trp Thr Gly Gly Ser Glu
 145 150 155 160
 His Phe Ser Met Ser Ser Leu His Tyr Thr Ala Phe Asp Pro Leu Phe
 165 170 175
 Tyr Phe His His Ser Asn Val Asp Arg Leu Trp Ala Val Trp Gln Ala
 180 185 190

Leu Gln Met Arg Arg His Lys Pro Tyr Arg Ala His Cys Ala Ile Ser
 195 200 205
 Leu Glu His Met His Leu Lys Pro Phe Ala Phe Ser Ser Pro Leu Asn
 210 215 220
 Asn Asn Glu Lys Thr His Ala Asn Ala Met Pro Asn Lys Ile Tyr Asp
 225 230 235 240
 Tyr Glu Asn Val Leu His Tyr Thr Tyr Glu Asp Leu Thr Phe Gly Gly
 245 250 255
 Ile Ser Leu Glu Asn Ile Glu Lys Met Ile His Glu Asn Gln Gln Glu
 260 265 270
 Asp Arg Ile Tyr Ala Gly Phe Leu Leu Ala Gly Ile Arg Thr Ser Ala
 275 280 285
 Asn Val Asp Ile Phe Ile Lys Thr Thr Asp Ser Val Gln His Lys Ala
 290 295 300
 Gly Thr Phe Ala Val Leu Gly Gly Ser Lys Glu Met Lys Trp Gly Phe
 305 310 315 320
 Asp Arg Val Phe Lys Phe Asp Ile Thr His Val Leu Lys Asp Leu Asp
 325 330 335
 Leu Thr Ala Asp Gly Asp Phe Glu Val Thr Val Asp Ile Thr Glu Val
 340 345 350
 Asp Gly Thr Lys Leu Ala Ser Ser Leu Ile Pro His Ala Ser Val Ile
 355 360 365
 Arg Glu His Ala Arg Gly Lys Leu Asn Arg
 370 375

<210> 70

<211> 419

<212> PRT

<213> Megathura crenulata

<400> 70

Asp Ser Ala His Thr Asp Asp Gly His Thr Glu Pro Val Met Ile Arg
 1 5 10 15
 Lys Asp Ile Thr Gln Leu Asp Lys Arg Gln Gln Leu Ser Leu Val Lys
 20 25 30
 Ala Leu Glu Ser Met Lys Ala Asp His Ser Ser Asp Gly Phe Gln Ala
 35 40 45
 Ile Ala Ser Phe His Ala Leu Pro Pro Leu Cys Pro Ser Pro Ala Ala
 50 55 60

Ser Lys Arg Phe Ala Cys Cys Val His Gly Met Ala Thr Phe Pro Gln
 65 70 75 80
 Trp His Arg Leu Tyr Thr Val Gln Phe Gln Asp Ser Leu Arg Lys His
 85 90 95
 Gly Ala Val Val Gly Leu Pro Tyr Trp Asp Trp Thr Leu Pro Arg Ser
 100 105 110
 Glu Leu Pro Glu Leu Leu Thr Val Ser Thr Ile His Asp Pro Glu Thr
 115 120 125
 Gly Arg Asp Ile Pro Asn Pro Phe Ile Gly Ser Lys Ile Glu Phe Glu
 130 135 140
 Gly Glu Asn Val His Thr Lys Arg Asp Ile Asn Arg Asp Arg Leu Phe
 145 150 155 160
 Gln Gly Ser Thr Lys Thr His His Asn Trp Phe Ile Glu Gln Ala Leu
 165 170 175
 Leu Ala Leu Glu Gln Thr Asn Tyr Cys Asp Phe Glu Val Gln Phe Glu
 180 185 190
 Ile Met His Asn Gly Val His Thr Trp Val Gly Gly Lys Glu Pro Tyr
 195 200 205
 Gly Ile Gly His Leu His Tyr Ala Ser Tyr Asp Pro Leu Phe Tyr Ile
 210 215 220
 His His Ser Gln Thr Asp Arg Ile Trp Ala Ile Trp Gln Ser Leu Gln
 225 230 235 240
 Arg Phe Arg Gly Leu Ser Gly Ser Glu Ala Asn Cys Ala Val Asn Leu
 245 250 255
 Met Lys Thr Pro Leu Lys Pro Phe Ser Phe Gly Ala Pro Tyr Asn Leu
 260 265 270
 Asn Asp His Thr His Asp Phe Ser Lys Pro Glu Asp Thr Phe Asp Tyr
 275 280 285
 Gln Lys Phe Gly Tyr Ile Tyr Asp Thr Leu Glu Phe Ala Gly Trp Ser
 290 295 300
 Ile Arg Gly Ile Asp His Ile Val Arg Asn Arg Gln Glu His Ser Arg
 305 310 315 320
 Val Phe Ala Gly Phe Leu Leu Glu Gly Phe Gly Thr Ser Ala Thr Val
 325 330 335
 Asp Phe Gln Val Cys Arg Thr Ala Gly Asp Cys Glu Asp Ala Gly Tyr
 340 345 350
 Phe Thr Val Leu Gly Gly Glu Lys Glu Met Pro Trp Ala Phe Asp Arg
 355 360 365

63

Leu Tyr Lys Tyr Asp Ile Thr Glu Thr Leu Asp Lys Met Asn Leu Arg
 370 375 380

His Asp Glu Ile Phe Gln Ile Glu Val Thr Ile Thr Ser Tyr Asp Gly
 385 390 395 400

Thr Val Leu Asp Ser Gly Leu Ile Pro Thr Pro Ser Ile Ile Tyr Asp
 405 410 415

Pro Ala His

<210> 71

<211> 418

<212> PRT

<213> Megathura crenulata

<400> 71

His Asp Ile Ser Ser His His Leu Ser Leu Asn Lys Val Arg His Asp
 1 5 10 15

Leu Ser Thr Leu Ser Glu Arg Asp Ile Gly Ser Leu Lys Tyr Ala Leu
 20 25 30

Ser Ser Leu Gln Ala Asp Thr Ser Ala Asp Gly Phe Ala Ala Ile Ala
 35 40 45

Ser Phe His Gly Leu Pro Ala Lys Cys Asn Asp Ser His Asn Asn Glu
 50 55 60

Val Ala Cys Cys Ile His Gly Met Pro Thr Phe Pro His Trp His Arg
 65 70 75 80

Leu Tyr Thr Leu Gln Phe Glu Gln Ala Leu Arg Arg His Gly Ser Ser
 85 90 95

Val Ala Val Pro Tyr Trp Asp Trp Thr Lys Pro Ile His Asn Ile Pro
 100 105 110

His Leu Phe Thr Asp Lys Glu Tyr Tyr Asp Val Trp Arg Asn Lys Val
 115 120 125

Met Pro Asn Pro Phe Ala Arg Gly Tyr Val Pro Ser His Asp Thr Tyr
 130 135 140

Thr Val Arg Asp Val Gln Glu Gly Leu Phe His Leu Thr Ser Thr Gly
 145 150 155 160

Glu His Ser Ala Leu Leu Asn Gln Ala Leu Leu Ala Leu Glu Gln His
 165 170 175

Asp Tyr Cys Asp Phe Ala Val Gln Phe Glu Val Met His Asn Thr Ile
 180 185 190

His Tyr Leu Val Gly Gly Pro Gln Val Tyr Ser Leu Ser Ser Leu His
 195 200 205

ERSATZBLATT (REGEL 26)

64

Tyr Ala Ser Tyr Asp Pro Ile Phe Phe Ile His His Ser Phe Val Asp
 210 215 220
 Lys Val Trp Ala Val Trp Gln Ala Leu Gln Glu Lys Arg Gly Leu Pro
 225 230 235 240
 Ser Asp Arg Ala Asp Cys Ala Val Ser Leu Met Thr Gln Asn Met Arg
 245 250 255
 Pro Phe His Tyr Glu Ile Asn His Asn Gln Phe Thr Lys Lys His Ala
 260 265 270
 Val Pro Asn Asp Val Phe Lys Tyr Glu Leu Leu Gly Tyr Arg Tyr Asp
 275 280 285
 Asn Leu Glu Ile Gly Gly Met Asn Leu His Glu Ile Glu Lys Glu Ile
 290 295 300
 Lys Asp Lys Gln His His Val Arg Val Phe Ala Gly Phe Leu Leu His
 305 310 315 320
 Gly Ile Arg Thr Ser Ala Asp Val Gln Phe Gln Ile Cys Lys Thr Ser
 325 330 335
 Glu Asp Cys His His Gly Gly Gln Ile Phe Val Leu Gly Gly Thr Lys
 340 345 350
 Glu Met Ala Trp Ala Tyr Asn Arg Leu Phe Lys Tyr Asp Ile Thr His
 355 360 365
 Ala Leu His Asp Ala His Ile Thr Pro Glu Asp Val Phe His Pro Ser
 370 375 380
 Glu Pro Phe Phe Ile Lys Val Ser Val Thr Ala Val Asn Gly Thr Val
 385 390 395 400
 Leu Pro Ala Ser Ile Leu His Ala Pro Thr Ile Ile Tyr Glu Pro Gly
 405 410 415
 Leu Gly

<210> 72
 <211> 241
 <212> PRT
 <213> Megathura crenulata

<400> 72
 Asp His His Glu Asp His His Ser Ser Ser Met Ala Gly His Gly Val
 1 5 10 15
 Arg Lys Glu Ile Asn Thr Leu Thr Thr Ala Glu Val Asp Asn Leu Lys
 20 25 30

65
 Asp Ala Met Arg Ala Val Met Ala Asp His Gly Pro Asn Gly Tyr Gln
 35 40 45
 Ala Ile Ala Ala Phe His Gly Asn Pro Pro Met Cys Pro Met Pro Asp
 50 55 60
 Gly Lys Asn Tyr Ser Cys Cys Thr His Gly Met Ala Thr Phe Pro His
 65 70 75 80
 Trp His Arg Leu Tyr Thr Lys Gln Met Glu Asp Ala Leu Thr Ala His
 85 90 95
 Gly Ala Arg Val Gly Leu Pro Tyr Trp Asp Gly Thr Thr Ala Phe Thr
 100 105 110
 Ala Leu Pro Thr Phe Val Thr Asp Glu Glu Asp Asn Pro Phe His His
 115 120 125
 Gly His Ile Asp Tyr Leu Gly Val Asp Thr Thr Arg Ser Pro Arg Asp
 130 135 140
 Lys Leu Phe Asn Asp Pro Glu Arg Gly Ser Glu Ser Phe Phe Tyr Arg
 145 150 155 160
 Gln Val Leu Leu Ala Leu Glu Gln Thr Asp Phe Cys Gln Phe Glu Val
 165 170 175
 Gln Phe Glu Ile Thr His Asn Ala Ile His Ser Trp Thr Gly Gly Leu
 180 185 190
 Thr Pro Tyr Gly Met Ser Thr Leu Glu Tyr Thr Thr Tyr Asp Pro Leu
 195 200 205
 Phe Trp Leu His His Ala Asn Thr Asp Arg Ile Trp Ala Ile Trp Gln
 210 215 220
 Ala Leu Gln Glu Tyr Arg Gly Leu Pro Tyr Asp His Ala Asn Cys Glu
 225 230 235 240
 Ile

<210> 73
 <211> 98
 <212> PRT
 <213> Megathura crenulata

<400> 73
 Lys His His Glu Lys His His Glu Asp His His Glu Asp Ile Leu Val
 1 5 10 15
 Arg Lys Asn Ile His Ser Leu Ser His His Glu Ala Glu Glu Leu Arg
 20 25 30
 Asp Ala Leu Tyr Lys Leu Gln Asn Asp Glu Ser His Gly Gly Tyr Glu
 35 40 45

66

His Ile Ala Gly Phe His Gly Tyr Pro Asn Leu Cys Pro Glu Lys Gly
 50 55 60
 Asp Glu Lys Tyr Pro Cys Cys Val His Gly Met Ser Ile Phe Pro His
 65 70 75 80
 Trp His Arg Leu His Thr Ile Gln Leu Glu Arg Ala Leu Lys Lys His
 85 90 95
 Gly Ser

<210> 74
 <211> 314
 <212> PRT
 <213> Megathura crenulata

<400> 74
 Gly Leu Pro Tyr Trp Asp Trp Thr Met Pro Met Ser His Leu Pro Glu
 1 5 10 15
 Leu Ala Thr Ser Glu Thr Tyr Leu Asp Pro Val Thr Gly Glu Thr Lys
 20 25 30
 Asn Asn Pro Phe His His Ala Gln Val Ala Phe Glu Asn Gly Val Thr
 35 40 45
 Ser Arg Asn Pro Asp Ala Lys Leu Phe Met Lys Pro Thr Tyr Gly Asp
 50 55 60
 His Thr Tyr Leu Phe Asp Ser Met Ile Tyr Ala Phe Glu Gln Glu Asp
 65 70 75 80
 Phe Cys Asp Phe Glu Val Gln Tyr Glu Leu Thr His Asn Ala Ile His
 85 90 95
 Ala Trp Val Gly Gly Ser Glu Lys Tyr Ser Met Ser Ser Leu His Tyr
 100 105 110
 Thr Ala Phe Asp Pro Ile Phe Tyr Leu His His Ser Asn Val Asp Arg
 115 120 125
 Leu Trp Ala Ile Trp Gln Ala Leu Gln Ile Arg Arg Gly Lys Ser Tyr
 130 135 140
 Lys Ala His Cys Ala Ser Ser Gln Glu Arg Glu Pro Leu Lys Pro Phe
 145 150 155 160
 Ala Phe Ser Ser Pro Leu Asn Asn Asn Glu Lys Thr Tyr His Asn Ser
 165 170 175
 Val Pro Thr Asn Val Tyr Asp Tyr Val Gly Val Leu His Tyr Arg Tyr
 180 185 190

67

Asp Asp Leu Gln Phe Gly Gly Met Thr Met Ser Glu Leu Glu Glu Tyr
 195 200 205

Ile His Lys Gln Thr Gln His Asp Arg Thr Phe Ala Gly Phe Phe Leu
 210 215 220

Ser Tyr Ile Gly Thr Ser Ala Ser Val Asp Ile Phe Ile Asn Arg Glu
 225 230 235 240

Gly His Asp Lys Tyr Lys Val Gly Ser Phe Val Val Leu Gly Gly Ser
 245 250 255

Lys Glu Met Lys Trp Gly Phe Asp Arg Met Tyr Lys Tyr Glu Ile Thr
 260 265 270

Glu Ala Leu Lys Thr Leu Asn Val Ala Val Asp Asp Gly Phe Ser Ile
 275 280 285

Thr Val Glu Ile Thr Asp Val Asp Gly Ser Pro Pro Ser Ala Asp Leu
 290 295 300

Ile Pro Pro Pro Ala Ile Ile Phe Glu Arg
 305 310

<210> 75
 <211> 416
 <212> PRT
 <213> Megathura crenulata

<400> 75

Ala Asp Ala Lys Asp Phe Gly His Ser Arg Lys Ile Arg Lys Ala Val
 1 5 10 15

Asp Ser Leu Thr Val Glu Glu Gln Thr Ser Leu Arg Arg Ala Met Ala
 20 25 30

Asp Leu Gln Asp Asp Lys Thr Ser Gly Gly Phe Gln Gln Ile Ala Ala
 35 40 45

Phe His Gly Glu Pro Lys Trp Cys Pro Ser Pro Glu Ala Glu Lys Lys
 50 55 60

Phe Ala Cys Cys Val His Gly Met Ala Val Phe Pro His Trp His Arg
 65 70 75 80

Leu Leu Thr Val Gln Gly Glu Asn Ala Leu Arg Lys His Gly Phe Thr
 85 90 95

Gly Gly Leu Pro Tyr Trp Asp Trp Thr Arg Ser Met Ser Ala Leu Pro
 100 105 110

His Phe Val Ala Asp Pro Thr Tyr Asn Asp Ala Ile Ser Ser Gln Glu
 115 120 125

Glu Asp Asn Pro Trp His His Gly His Ile Asp Ser Val Gly His Asp
 130 135 140

68

Thr Thr Arg Asp Val Arg Asp Asp Leu Tyr Gln Ser Pro Gly Phe Gly
 145 150 155 160
 His Tyr Thr Asp Ile Ala Lys Gln Val Leu Leu Ala Phe Glu Gln Asp
 165 170 175
 Asp Phe Cys Asp Phe Glu Val Gln Phe Glu Ile Ala His Asn Phe Ile
 180 185 190
 His Ala Leu Val Gly Gly Asn Glu Pro Tyr Ser Met Ser Ser Leu Arg
 195 200 205
 Tyr Thr Thr Tyr Asp Pro Ile Phe Phe Leu His Arg Ser Asn Thr Asp
 210 215 220
 Arg Leu Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Lys Pro
 225 230 235 240
 Tyr Asn Thr Ala Asn Cys Ala Ile Ala Ser Met Arg Lys Pro Leu Gln
 245 250 255
 Pro Phe Gly Leu Asp Ser Val Ile Asn Pro Asp Asp Glu Thr Arg Glu
 260 265 270
 His Ser Val Pro Phe Arg Val Phe Asp Tyr Lys Asn Asn Phe Asp Tyr
 275 280 285
 Glu Tyr Glu Ser Leu Ala Phe Asn Gly Leu Ser Ile Ala Gln Leu Asp
 290 295 300
 Arg Glu Leu Gln Arg Arg Lys Ser His Asp Arg Val Phe Ala Gly Phe
 305 310 315 320
 Leu Leu His Glu Ile Gly Gln Ser Ala Leu Val Lys Phe Tyr Val Cys
 325 330 335
 Lys His Asn Val Ser Asp Cys Asp His Tyr Ala Gly Glu Phe Tyr Ile
 340 345 350
 Leu Gly Asp Glu Ala Glu Met Pro Trp Arg Tyr Asp Arg Val Tyr Lys
 355 360 365
 Tyr Glu Ile Thr Gln Gln Leu His Asp Leu Asp Leu His Val Gly Asp
 370 375 380
 Asn Phe Phe Leu Lys Tyr Glu Ala Phe Asp Leu Asn Gly Gly Ser Leu
 385 390 395 400
 Gly Gly Ser Ile Phe Ser Gln Pro Ser Val Ile Phe Glu Pro Ala Ala
 405 410 415

<210> 76

<211> 419

<212> PRT

69

<213> Megathura crenulata

<400> 76

Gly Ser His Gln Ala Asp Glu Tyr Arg Glu Ala Val Thr Ser Ala Ser
 1 5 10 15
 His Ile Arg Lys Asn Ile Arg Asp Leu Ser Glu Gly Glu Ile Glu Ser
 20 25 30
 Ile Arg Ser Ala Phe Leu Gln Ile Gln Lys Glu Gly Ile Tyr Glu Asn
 35 40 45
 Ile Ala Lys Phe His Gly Lys Pro Gly Leu Cys Glu His Asp Gly His
 50 55 60
 Pro Val Ala Cys Cys Val His Gly Met Pro Thr Phe Pro His Trp His
 65 70 75 80
 Arg Leu Tyr Val Leu Gln Val Glu Asn Ala Leu Leu Glu Arg Gly Ser
 85 90 95
 Ala Val Ala Val Pro Tyr Trp Asp Trp Thr Glu Lys Ala Asp Ser Leu
 100 105 110
 Pro Ser Leu Ile Asn Asp Ala Thr Tyr Phe Asn Ser Arg Ser Gln Thr
 115 120 125
 Phe Asp Pro Asn Pro Phe Phe Arg Gly His Ile Ala Phe Glu Asn Ala
 130 135 140
 Val Thr Ser Arg Asp Pro Gln Pro Glu Leu Trp Asp Asn Lys Asp Phe
 145 150 155 160
 Tyr Glu Asn Val Met Leu Ala Leu Glu Gln Asp Asn Phe Cys Asp Phe
 165 170 175
 Glu Ile Gln Leu Glu Leu Ile His Asn Ala Leu His Ser Arg Leu Gly
 180 185 190
 Gly Arg Ala Lys Tyr Ser Leu Ser Ser Leu Asp Tyr Thr Ala Phe Asp
 195 200 205
 Pro Val Phe Phe Leu His His Ala Asn Val Asp Arg Ile Trp Ala Ile
 210 215 220
 Trp Gln Asp Leu Gln Arg Tyr Arg Lys Lys Pro Tyr Asn Glu Ala Asp
 225 230 235 240
 Cys Ala Val Asn Glu Met Arg Lys Pro Leu Gln Pro Phe Asn Asn Pro
 245 250 255
 Glu Leu Asn Ser Asp Ser Met Thr Leu Lys His Asn Leu Pro Gln Asp
 260 265 270
 Ser Phe Asp Tyr Gln Asn Arg Phe Arg Tyr Gln Tyr Asp Asn Leu Gln
 275 280 285

70

Phe Asn His Phe Ser Ile Gln Lys Leu Asp Gln Thr Ile Gln Ala Arg
 290 295 300
 Lys Gln His Asp Arg Val Phe Ala Gly Phe Ile Leu His Asn Ile Gly
 305 310 315 320
 Thr Ser Ala Val Val Asp Ile Tyr Ile Cys Val Glu Gln Gly Gly Glu
 325 330 335
 Gln Asn Cys Lys Thr Lys Ala Gly Ser Phe Thr Ile Leu Gly Gly Glu
 340 345 350
 Thr Glu Met Pro Phe His Phe Asp Arg Leu Tyr Lys Phe Asp Ile Thr
 355 360 365
 Ser Ala Leu His Lys Leu Gly Val Pro Leu Asp Gly His Gly Phe Asp
 370 375 380
 Ile Lys Val Asp Val Arg Ala Val Asn Gly Ser His Leu Asp Gln His
 385 390 395 400
 Ile Leu Asn Glu Pro Ser Leu Leu Phe Val Pro Gly Glu Arg Lys Asn
 405 410 415
 Ile Tyr Tyr

<210> 77

<211> 413

<212> PRT

<213> Megathura crenulata

<400> 77

Asp Gly Leu Ser Gln His Asn Leu Val Arg Lys Glu Val Ser Ser Leu
 1 5 10 15
 Thr Thr Leu Glu Lys His Phe Leu Arg Lys Ala Leu Lys Asn Met Gln
 20 25 30
 Ala Asp Asp Ser Pro Asp Gly Tyr Gln Ala Ile Ala Ser Phe His Ala
 35 40 45
 Leu Pro Pro Leu Cys Pro Ser Pro Ser Ala Ala His Arg His Ala Cys
 50 55 60
 Cys Leu His Gly Met Ala Thr Phe Pro Gln Trp His Arg Leu Tyr Thr
 65 70 75 80
 Val Gln Phe Glu Asp Ser Leu Lys Arg His Gly Ser Ile Val Gly Leu
 85 90 95
 Pro Tyr Trp Asp Trp Leu Lys Pro Gln Ser Ala Leu Pro Asp Leu Val
 100 105 110
 Thr Gln Glu Thr Tyr Glu His Leu Phe Ser His Lys Thr Phe Pro Asn
 115 120 125

Pro Phe Leu Lys Ala Asn Ile Glu Phe Glu Gly Glu Gly Val Thr Thr
 130 135 140
 Glu Arg Asp Val Asp Ala Glu His Leu Phe Ala Lys Gly Asn Leu Val
 145 150 155 160
 Tyr Asn Asn Trp Phe Cys Asn Gln Ala Leu Tyr Ala Leu Glu Gln Glu
 165 170 175
 Asn Tyr Cys Asp Phe Glu Ile Gln Phe Glu Ile Leu His Asn Gly Ile
 180 185 190
 His Ser Trp Val Gly Gly Ser Lys Thr His Ser Ile Gly His Leu His
 195 200 205
 Tyr Ala Ser Tyr Asp Pro Leu Phe Tyr Ile His His Ser Gln Thr Asp
 210 215 220
 Arg Ile Trp Ala Ile Trp Gln Ala Leu Gln Glu His Arg Gly Leu Ser
 225 230 235 240
 Gly Lys Glu Ala His Cys Ala Leu Glu Gln Met Lys Asp Pro Leu Lys
 245 250 255
 Pro Phe Ser Phe Gly Ser Pro Tyr Asn Leu Asn Lys Arg Thr Gln Glu
 260 265 270
 Phe Ser Lys Pro Glu Asp Thr Phe Asp Tyr His Arg Phe Gly Tyr Glu
 275 280 285
 Tyr Asp Ser Leu Glu Phe Val Gly Met Ser Val Ser Ser Leu His Asn
 290 295 300
 Tyr Ile Lys Gln Gln Gln Glu Ala Asp Arg Val Phe Ala Gly Phe Leu
 305 310 315 320
 Leu Lys Gly Phe Gly Gln Ser Ala Ser Val Ser Phe Asp Ile Cys Arg
 325 330 335
 Pro Asp Gln Ser Cys Gln Glu Ala Gly Tyr Phe Ser Val Leu Gly Gly
 340 345 350
 Ser Ser Glu Met Pro Trp Gln Phe Asp Arg Leu Tyr Lys Tyr Asp Ile
 355 360 365
 Thr Lys Thr Leu Lys Asp Met Lys Leu Arg Tyr Asp Asp Thr Phe Thr
 370 375 380
 Ile Lys Val His Ile Lys Asp Ile Ala Gly Ala Glu Leu Asp Ser Asp
 385 390 395 400
 Leu Ile Pro Thr Pro Ser Val Leu Leu Glu Gly Lys
 405 410

72

<210> 78

<211> 417

<212> PRT

<213> Megathura crenulata

<400> 78

His Gly Ile Asn Val Arg His Val Gly Arg Asn Arg Ile Arg Met Glu
 1 5 10 15
 Leu Ser Glu Leu Thr Glu Arg Asp Leu Ala Ser Leu Lys Ser Ala Met
 20 25 30
 Arg Ser Leu Gln Ala Asp Asp Gly Val Asn Gly Tyr Gln Ala Ile Ala
 35 40 45
 Ser Phe His Gly Leu Pro Ala Ser Cys His Asp Asp Glu Gly His Glu
 50 55 60
 Ile Ala Cys Cys Ile His Gly Met Pro Val Phe Pro His Trp His Arg
 65 70 75 80
 Leu Tyr Thr Leu Gln Met Asp Met Ala Leu Leu Ser His Gly Ser Ala
 85 90 95
 Val Ala Ile Pro Tyr Trp Asp Trp Thr Lys Pro Ile Ser Lys Leu Pro
 100 105 110
 Asp Leu Phe Thr Ser Pro Glu Tyr Tyr Asp Pro Trp Arg Asp Ala Val
 115 120 125
 Val Asn Asn Pro Phe Ala Lys Gly Tyr Ile Lys Ser Glu Asp Ala Tyr
 130 135 140
 Thr Val Arg Asp Pro Gln Asp Ile Leu Tyr His Leu Gln Asp Glu Thr
 145 150 155 160
 Gly Thr Ser Val Leu Leu Asp Gln Thr Leu Leu Ala Leu Glu Gln Thr
 165 170 175
 Asp Phe Cys Asp Phe Glu Val Gln Phe Glu Val Val His Asn Ala Ile
 180 185 190
 His Tyr Leu Val Gly Gly Arg Gln Val Tyr Ala Leu Ser Ser Gln His
 195 200 205
 Tyr Ala Ser Tyr Asp Pro Ala Phe Phe Ile His His Ser Phe Val Asp
 210 215 220
 Lys Ile Trp Ala Val Trp Gln Ala Leu Gln Lys Lys Arg Lys Arg Pro
 225 230 235 240
 Tyr His Lys Ala Asp Cys Ala Leu Asn Met Met Thr Lys Pro Met Arg
 245 250 255
 Pro Phe Ala His Asp Phe Asn His Asn Gly Phe Thr Lys Met His Ala
 260 265 270

73

Val Pro Asn Thr Leu Phe Asp Phe Gln Asp Leu Phe Tyr Thr Tyr Asp
 275 280 285

Asn Leu Glu Ile Ala Gly Met Asn Val Asn Gln Leu Glu Ala Glu Ile
 290 295 300

Asn Arg Arg Lys Ser Gln Thr Arg Val Phe Ala Gly Phe Leu Leu His
 305 310 315 320

Gly Ile Gly Arg Ser Ala Asp Val Arg Phe Trp Ile Cys Lys Thr Ala
 325 330 335

Asp Asp Cys His Ala Ser Gly Met Ile Phe Ile Leu Gly Gly Ser Lys
 340 345 350

Glu Met His Trp Ala Tyr Asp Arg Asn Phe Lys Tyr Asp Ile Thr Gln
 355 360 365

Ala Leu Lys Ala Gln Ser Ile His Pro Glu Asp Val Phe Asp Thr Asp
 370 375 380

Ala Pro Phe Phe Ile Lys Val Glu Val His Gly Val Asn Lys Thr Ala
 385 390 395 400

Leu Pro Ser Ser Ala Ile Pro Ala Pro Thr Ile Ile Tyr Ser Ala Gly
 405 410 415

Glu

<210> 79
 <211> 395
 <212> PRT
 <213> Megathura crenulata

<400> 79
 Asp His Ile Ala Gly Ser Gly Val Arg Lys Asp Val Thr Ser Leu Thr
 1 5 10 15

Ala Ser Glu Ile Glu Asn Leu Arg His Ala Leu Gln Ser Val Met Asp
 20 25 30

Asp Asp Gly Pro Asn Gly Phe Gln Ala Ile Ala Ala Tyr His Gly Ser
 35 40 45

Pro Pro Met Cys His Met Xaa Asp Gly Arg Asp Val Ala Cys Cys Thr
 50 55 60

His Gly Met Ala Ser Phe Pro His Trp His Arg Leu Phe Val Lys Gln
 65 70 75 80

Met Glu Asp Ala Leu Ala Ala His Gly Ala His Ile Gly Ile Pro Tyr
 85 90 95

Trp Asp Trp Thr Ser Ala Phe Ser His Leu Pro Ala Leu Val Thr Asp
 100 105 110

74

His Glu His Asn Pro Phe His His Gly His Ile Ala His Arg Asn Val
 115 120 125
 Asp Thr Ser Arg Ser Pro Arg Asp Met Leu Phe Asn Asp Pro Glu His
 130 135 140
 Gly Ser Glu Ser Phe Phe Tyr Arg Gln Val Leu Leu Ala Leu Glu Gln
 145 150 155 160
 Thr Asp Phe Cys Gln Phe Glu Val Gln Phe Glu Ile Thr His Asn Ala
 165 170 175
 Ile His Ser Trp Thr Gly Gly His Thr Pro Tyr Gly Met Ser Ser Leu
 180 185 190
 Glu Tyr Thr Ala Tyr Asp Pro Leu Phe Tyr Leu His His Ser Asn Thr
 195 200 205
 Asp Arg Ile Trp Ala Ile Trp Gln Ala Leu Gln Lys Tyr Arg Gly Phe
 210 215 220
 Gln Tyr Asn Ala Ala His Cys Asp Ile Gln Val Leu Lys Gln Pro Leu
 225 230 235 240
 Lys Pro Phe Ser Glu Ser Arg Asn Pro Asn Pro Val Thr Arg Ala Asn
 245 250 255
 Ser Arg Ala Val Asp Ser Phe Asp Tyr Glu Arg Leu Asn Tyr Gln Tyr
 260 265 270
 Asp Thr Leu Thr Phe His Gly His Ser Ile Ser Glu Leu Asp Ala Met
 275 280 285
 Leu Gln Glu Arg Lys Lys Glu Glu Arg Thr Phe Ala Ala Phe Leu Leu
 290 295 300
 His Gly Phe Gly Ala Ser Ala Asp Val Ser Phe Asp Val Cys Thr Pro
 305 310 315 320
 Asp Gly His Cys Ala Phe Ala Gly Thr Phe Ala Val Leu Gly Gly Glu
 325 330 335
 Leu Glu Met Pro Trp Ser Phe Glu Arg Leu Phe Arg Tyr Asp Ile Thr
 340 345 350
 Lys Val Leu Lys Gln Met Asn Leu His Tyr Asp Ser Glu Phe His Phe
 355 360 365
 Glu Leu Lys Ile Val Gly Thr Asp Gly Thr Glu Leu Pro Ser Asp Arg
 370 375 380
 Ile Lys Ser Pro Thr Ile Glu His His Gly Gly
 385 390 395

<210> 80
 <211> 1266
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 80
 cttgttcagtt ttctactcgt cgccttctgt gtgggggctg gagcagacaa cgtcgtcaga 60
 aaggacgtga gtcacctcac ggatgacgag gtgcaagctc tccacggcgc cctccatgac 120
 gtcactgcat ctacagggcc tctgagtttc gaagacataa catcttacca tgccgcacca 180
 gcgtcgtgtg actacaaggg acggaagatc gcctgctgtg tccacggtat gccagtttc 240
 cccttctggc acagggcata tgctgtccaa gccgagcggg cactgttgctc caaacggaag 300
 actgtcggaa tgccttactg ggactggacg caaacgctga ctacttacc atctcttggtg 360
 actgaaccca tctacattga cagtaaaggt ggaaaggctc aaaccaacta ctggtagcgc 420
 ggcgagatag cgttcatcaa taagaagact gcgcgagctg tagatgatcg cctattcgag 480
 aagggtggagc ctgggtcacta cacacatctt atggagactg tcctcgacgc tctcgaacag 540
 gacgaattct gtaaatttga aatccagttc gagttggctc ataattgctat ccattacttg 600
 gttggcggta aatttgaata ttcaatgtca aacttggaaat acacctccta cgaccccatc 660
 ttcttctctc accactccaa cgttgaccgc ctcttcgcca tctggcagcg tcttcaggaa 720
 ctgagaggaa agaattccaa tgcaatggac tgtgcacatg aactcgctca ccagcaactc 780
 caaccttca acagggacag caatccagtc cagctcacia aggaccactc gacacctgct 840
 gacctctttg attacaaaca acttggtatc agctacgaca gcttaaacct gaattggaatg 900
 acgccagaac agctgaaaac agaactagac gaacgccact ccaaagaacg tgcgtttgca 960
 agcttccgac tcagtggctt tgggggttct gccaacgttg ttgtctatgc atgtgtccct 1020
 gatgatgatc cagcagtgat tgactactgc gagaaagcag gcgacttctt cattcttggg 1080
 ggtcaaagcg aaatgccgtg gagattctac agacccttct tctatgatgt aactgaagcg 1140
 gtacatcacc ttggagtccc gctaagtggc cactactatg tgaaaacaga actcttcagc 1200
 gtgaatggca cagcactttc acctgatctt ctctctcaac caactgttgc ctaccgacct 1260
 ggga 1266

<210> 81
 <211> 1257
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 81
 ggtcaccttg accacactgt gcatcatcgc cagcatgacg atcttattgt tcgaaaaaat 60
 atagatcatt tgactcgtga agaggaatac gagctaagga tggctctgga gagattccag 120
 gccgacacat ccgttgatgg gtaccaggct acagtagagt accatggcct tccgtcgtgt 180
 tgtccacgac cagatgcaaa agtcaggttc gcctgttgta tgcatggcat ggcatccttc 240
 cctcactggc accggctgtt cgttaccacg gtggaagatg ctcttgtaag gcgtggatcg 300
 cctatcgggt ttccttattg ggactggaca aaacctatga ctacacctcc agacttggca 360
 tcaaattgaga cgtacgtaga cccttatgga catacacatc ataattccatt cttcaatgca 420
 aatatatctt ttgaggaggg acaccatcac acgagcagga tgatagattc gaaactgttt 480
 gccccagtcg cttttgggga gcattcccat ctgtttgatg gaatcctgta cgcatttgag 540
 caggaagatt tctgcgactt tgagattcag tttgagttag tccataattc tattcatgag 600
 tggataggcg gttccgaaga ttactccatg gccacctgac attacacagc ctttgacccc 660
 attttctacc ttcattcatt caatgtcgat cgtctatggg caatctggca agctcttcaa 720
 atcaggagac acaagccata tcaagccac tgtgcacagt ctgtggaaca gttgccaatg 780
 aagccatttg ctttcccatc acctcttaac aacaacgaga agacacatag tcattcagtc 840
 ccgactgaca tttatgacta cgagggaagt ctgactaca gctacgatga tctaactgtt 900
 ggtgggatga accttgaaga aatagaagaa ctatacatc tcagacaaca gcatgaacga 960
 gtcttcgagg gatttctcct tgctggaata ggaacatctg cacttgttga cattttcata 1020
 aataaaccgg ggaaccaacc actcaaagct ggagatattg ccattcttgg tgggtgcaaag 1080
 gaaatgcctt gggcgtttga ccgcttggat aaggctgaaa taactgactc attgaagaca 1140
 ctttctctcg atgtcgatgg agattatgaa gtcactttta aaattcatga tatgcacgga 1200
 aacgctcttg atacggacct gattccacac gcagcagttg tttctgagcc agctcac 1257

<210> 82
 <211> 1242
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 82
 cctacctttg aggatgaaaa gcacagctta cgaatcagaa aaaatgtcga cagcttgact 60
 cctgaagaaa caaatgaact gcgtaaagcc ctggagcttc ttgaaaatga tcatactgca 120
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 tatgatcacc tcttcattcg ctacgaagtc tttgatctta aaggagttag tttgggaact 1200
 gacctgttcc acactgcaaa tgtggtacat gattccggca ca 1242

<210> 83
 <211> 1239
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 83
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 caggacgacg gaacatatga atctattggc cagtaccatg gcaaaccagg caaatgtcaa 180
 ttgaatgacg ataattattg gtgtgtgtgc catggtatgc ctaccttccc ccagtggcac 240
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 gaccgtttgt ataaatttga aatcaacaaa ccactgcaac agttaggagt caagctgcat 1140
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 catacctttg atccaactat catctttgaa cctggaaca 1239

<210> 84
 <211> 1260
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 84
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 cgctccgctg atgggtacca agccattgcc tctttccatg ccctgccacc actctgtccc 180
 aatccatctg cagctcaccg ttatgcttgc tgtgtccatg gcatggctac atttccccag 240
 tggcacagac tgtacactgt tcagggttcag gatgccctga ggagacatgg ttcacttggt 300
 ggtattcctt actgggactg gacaaaacca gtcaacgagt taccgagct tctttcttca 360
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 atagaatttg aaggaccggg cgttcataca gagaggcaca taaatactga gcgcctgttt 480
 cacagtgggg atcatgacgg ataccacaac tggttcttcg aaactgttct ctttgctttg 540
 gaacaggaag attactgcga ttttgaaata caatttgaga tagcccataa tggcatccac 600
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 ccaattttct acatccacca ttcacagacg gacagaatat gggctatttg gcaagagctg 720
 cagaagtaca ggggtctatc tggttcggaa gcaaactgtg ccattgaaca tatgagaaca 780
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 gccgaaatgc catgggcatt cgacaggctt tataagtatg acattactaa aactcttcac 1140
 gacatgaacc tgaggcacga ggacacttct tctatagacy taactatcac gtcttacaat 1200
 ggaacagtac tctcgggaga cctcattcag acgcctcca ttatatttgt acctggacgc 1260

<210> 85
 <211> 1251
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 85
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 agttcccgtg acatagcaag cttgaaggca gctttgacaa gccttcaaca tgataatggg 120
 actgatgggt atcaagctat tgctgccttc catggcgctt ctgcgcagtg ccacgagcca 180
 tctggacgtg agatcgctg ttgcatccac ggcatggcga cgtttctca ctggcaccgg 240
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<210> 86
 <211> 1209
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 86
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 gaccacggtc ccaatggctt tcaagctatt gctgctttcc atggaaaacc agcttttgtt 180
 cccatgcctg atggccacaa ctactcatgt tgtactcacg gcatggctac cttcccacac 240
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 ggcctgacct actgggactg gactgctgcc ttcacccacc tgccaacact ggtcaccgac 360
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 gaaccgggc 1209

<210> 87
 <211> 1536
 <212> DNA
 <213> *Haliotis tuberculata*

<400> 87
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 atcaggaaaag aagttgactt cctctccctg caagaggcca acgcaattaa ggatgcactg 120
 tacaagctcc agaatgacga cagtaaaggg ggctttgagg ccatagctgg ctatcacggg 180
 tatcctaata tgtgtccaga aagagggtacc gacaagtatc cctgctgtgt ccacggaatg 240
 cccgtgttcc cccactggca ccgacctgcat accattcaga tggagagagc tctgaaaaac 300
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 tctttctttg gagattccag caacaacaac cctttctaca aatattacat ccggggcggtg 420
 cagcacgaaa caaccaggga cattaatcag agactcttta atcaaaccac gtttggtgaa 480
 tttgattacc tatattacct aactctgcaa gtccctggagg aaaactcgta ctgtgacttt 540
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 gagtttacac caatagattc gtcggtgaac aaagcaatgg tggagctggg cagctatact 1380

79

gctatggcta aatgcatcgt tccccctttc tcttaccacg gctttgaact ggacaaagtc 1440
 tacagcgtcg atcacggaga ctactacatt gctgcaggta cccacgcgtt gtgtgagcag 1500
 aacctcaggc tccacatcca cgtggaacac gagtag 1536

<210> 88

<211> 591

<212> DNA

<213> *Haliotis tuberculata*

<400> 88

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 atcaagtttg agaataagaa gactgcaaga gctgttgacg atcgctttt cgagaaggtt 180
 ggaccaggag agaatacccg actctttgaa ggaattctcg atgctcttga acaggatgaa 240
 ttctgcaact tcgagatcca gtttgagttg gctcacaacg ctatccacta cctggttggc 300
 ggccgtcaca cgtactccat gtctcatctc gagtacacct cctacgacct cctcttcttc 360
 ctccatcact ccaacccgga ccgcatcttc gccatctggg aacgtcttca ggtactcaga 420
 ggaaaggacc ccaacaccgc cgactgcgca cacaacctca tccatgagcc catggaaccg 480
 ttccgtcggc atgagcccat ggaaccgttc cgtcgggact cgaaccctct tgacctcacc 540
 agggaaaact ccaaaccaat tgacagcttt gattatgcc ccttggtta c 591

<210> 89

<211> 1245

<212> DNA

<213> *Haliotis tuberculata*

<400> 89

gttacagagg cccagctcc ctctcggat gctcacctcg ccgtcaggaa ggatatcaac 60
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 ttctgcgatt ttgaagttca atttgagatg acccataata atattcacgc ctggattggt 600
 ggcggcgaga agtattccat gtcttctcta cactacacag ccttcgacct tatcttctac 660
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<210> 90

<211> 1251

<212> DNA

<213> *Haliotis assimilis*

<400> 90

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80

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ggtggattcc agcaaattgc tgcttttcac ggggaaccca aatggtgccc aagtcctcat 180
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gatacagttg gtgttgacac aacaagaagc gtccgtcaag aactgtatga agctcctgga 480
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<210> 91

<211> 1242

<212> DNA

<213> *Haliotis tuberculata*

<400> 91

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cagaacgacg gagtctatga gaatattgcc aagttccacg gcaagcctgg gttgtgtgat 180
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<210> 92

<211> 1257

<212> DNA

<213> *Haliotis tuberculata*

<400> 92

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81

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tcaccatctg cagctcaccg ttatgcttgc tgtgtccacg gcatggctac atttcccag 240
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actgtcctga gcagcagctt aatcccaaca ccgagtgtca tattccagcg gggacat 1257

```

<210> 93

<211> 1248

<212> DNA

<213> *Haliotis tuberculata*

<400> 93

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cgtgacataa ataccaggag catgtcaccg aaccgtgttc gccgtgagct gagcgatctg 60
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cccaacggat accaggctct tgcaaccttc catgggctac cagcaggctg ccatgatagc 180
cggggaaatg agatcgcatg ttgcattcac gggatgccga ccttcccca gtggcacaga 240
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tatgacccat ggcagatgac tgtggtaaac aaccattct ccaaagggtt tgtcaaattt 420
gcaaatacct acacagtaag agaccacag gagatgctgt tccagctttg tgaacatgga 480
gagtcacatc tctatgagca aactcttctt gctcttgagc aaaccgacta ctgtgatttt 540
gaggtagagt ttgaggtcct ccataacgtg atccactacc ttgttgggtg acgtcagacc 600
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gatgctgagg agccatttta tatcaagggt gagatccatg ctgttaacaa gaccatgata 1200
ccgtcgtctg tgatcccagc ccaactatc atctattctc ctggggaa 1248

```

<210> 94

<211> 1206

<212> DNA

<213> *Haliotis tuberculata*

<400> 94

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ggtcgcgctg ctgacagtgc gcactctgcc aacattgctg gctctggggt gaggaaggac 60
gtcacgaccc tcaactgtgtc tgagaccgag aacctaaagc aggtcttca aggtgtcatc 120
gatgatactg gtcccaatgg ttaccaagca atagcatcct tccacggaag tccccaatg 180
tgcgagatga acggccgcaa ggttgccgtg tgtgctcacg gtatggcctc cttcccacac 240

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82

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tggcacagac tgtatgtgaa gcagatggaa gatgccctgg ctgaccacgg gtcacatatc 300
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tccgagaaca atcccttcca tgagggtcgc attgatcatc tcggtgtaac cacgtcacgt 420
tccccagag acatgctgtt taacgaccca gagcaaggat cagagtcgtt cttctataga 480
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gctgtctggc aagcactgca gaaataccga ggactcccat acaacgaagc aactgtgaa 720
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agagtcatga atcagctcca tctccagtat gattcagatt tcagtttcag ggtgaagctt 1140
gttccacca atggcactga gctttcatca gaccttctca agtcaccaac aattgaacat 1200
gaactt                                     1206

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<210> 95

<211> 1548

<212> DNA

<213> *Haliotis tuberculata*

<400> 95

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ggagcccaca gaggaccagt tgaagaaaca gaagtcactc gccaacatac tgacggcaat 60
gcacactttc atcgtaagga agttgattcg ctgtccctgg atgaagcaaa caacttgaag 120
aatgcccttt acaagctaca gaacgaccac agtctaacgg gatacgaagc aatctctggg 180
taccatggat accccaatct gtgtccggaa gaaggcgatg acaaaatacc cctgctcggt 240
ccccggatgg gcacttttcc ttactggcac agactcttga ccattcaact ggaaagagct 300
cttgagcaca atggtgcaat gcttggtgtt cttactggg actggaacaa ggacctgtcg 360
tactgcccgg cgttcttctc cgactccagc aacaacaatc cctacttcaa gtaccacatc 420
gccggtgttg gtcacgacac cgtcagagag ccaactagtc ttatatataa ccagcccaa 480
atccatggtt atgattatct ctattaccta gcattgacca cgcttgaaga aaacaattac 540
tgggactttg aggttcagta tgagatcctc cacaacggcg tccactcctg gcttgaggga 600
tccagaagat attccatgtc taccctggag tattcgccct ttgacctgt ctttatgatc 660
cttactcggg gtctagacag actttggatc atctggcaag aacttcagaa gatcaggaga 720
aagccctaca acttcgctaa atgtgcttat catatgatgg aagagccact ggcgcccttc 780
agctatccat ctatcaacca ggacgagttc acccgtgcc aactccaaggc ttctacagtt 840
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caagagccta cccaaatcct ttcatcatct acagacctgc caatcatgac tacgatgttc 1260
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aaccacgtct attctgtcaa gcctggtgac tactatgtta ctggaccac gagagacctt 1500
tgccagaatg cagatgtcag gattcatatc catgttgagg atgagtaa 1548

```

<210> 96

<211> 966

<212> DNA

<213> *Megathura crenulata*

<400> 96

83

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ggcctaccgt actgggactg gactgaaccc atgacacaca ttccgggtct ggcaggaaac 60
aaaacttatg tggattctca tgggtgcatcc cacacaaatc cttttcatag ttcagtgtatt 120
gcatttgaag aaaatgctcc ccacacccaaa agacaaatag atcaaagact ctttaaacc 180
gctacctttg gacaccacac agacctgttc aaccagattt tgtatgcctt tgaacaagaa 240
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cggcataaac cctacagggc ccactgcgcc atatctctgg aacatatgca tctgaaacca 480
ttcgcccttt catctcccct taacaataac gaaaagactc atgccaatgc catgccaaac 540
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atctctctgg aaaacataga aaagatgata cacgaaaacc agcaagaaga cagaatatat 660
gccggttttc tcctggctgg catagctact tcagcaaagc ttgatattct cattaataat 720
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ctcactgctg atggcgattt cgaagttact gttgacatca ctgaagtcga tggaaactaa 900
cttgcataca gtcttattcc acatgcttct gtcattcgtg agcatgcacg tggtaagctg 966
aataga

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<210> 97

<211> 1242

<212> DNA

<213> Megathura crenulata

<400> 97

```

gttaaatttg acaaagtgcc aaggagtctg cttattcgaa aaaatgtaga ccgtttgagc 60
cccaggagaa tgaatgaact tcgtaaaagcc cttagccttac tgaaagagga caaaagtggc 120
ggtggatttc agcagcttgg tgcattccat ggggagccaa aatgggtgtc tagtcccgaa 180
gcatctaaaa aatttgcttg ctgtgttcac ggcatgtctg tgttccctca ctggcatcga 240
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tactgggatt ggacctctcc tcttaatcac cttcccgaac tggcagatca tgagaaqtac 360
gtcgaccctg aagatggggt agagaagcat aacccttggc tcgatgggtca tatagataca 420
gtcgacaaaa caacaacaag aagtgttcag aataaactct tcgaacagcc tgagtttggc 480
cattatacaa gcattgcaa acaagtactg ctagcgttgg aacaggacaa tttctgtgac 540
tttgaaatcc aatatgagat tgcccataac tacatccatg cacttgtagg aggcgctcag 600
ccttatggta tggcatcgct tcgctacact gcttttgatc cactattcta cttgcatcac 660
tctaatacag atcgatatg ggcaatatgg caggctttac agaagtacag aggaaaaccg 720
tacaacgttg ctaactgtgc tgttacatcg atgagagaac ctttgcaacc atttggcctc 780
tctgccaata tcaacacaga ccatgtaacc aaggagcatt cagtgcatt caacgttttt 840
gattacaaga ccaatttcaa ttatgaatat gacactttgg aatttaacgg tctctcaatc 900
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aactgtcacc ccgctggaga gttttacctt ctgggtgatg aaaacgagat gccatgggca 1080
tacgatagag tcttcaata tgacataacc gaaaaactcc acgatctaaa gctgcatgca 1140
gaagaccact tctacattga ctatgaagta tttgacctta aaccagcaag cctgggaaaa 1200
gatttgttca agcagccttc agtcattcat gaaccaagaa ta 1242

```

<210> 98

<211> 1236

<212> DNA

<213> Megathura crenulata

<400> 98

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ggtcaccatg aaggcgaagt atatcaagct gaagtaactt ctgccaaccg tattcgaaaa 60
aacattgaaa atctgagcct tgggtgaactc gaaagtctga gagctgcctt cctggaaatt 120
gaaaacgatg gaacttacga atcaatagct aaattccatg gtagccctgg tttgtgccag 180
ttaaatggta accccatctc ttgttgtgtc catggcatgc caactttccc tctctggcac 240
agactgtacg tggttgtcgt tgagaatgcc ctcctgaaaa aaggatcatc tgtagctgtt 300

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84

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ccctattggg actggacaaa acgaatcgaa catttacctc acctgatttc agacgccact 360
tactacaatt ccaggcaaca tcactatgag acaaaccat tccatcatgg caaaatcaca 420
cacgagaatg aaatcactac tagggatccc aaggacagcc tcttccattc agactacttt 480
tacgagcagg tccttttacgc cttggagcag gataacttct gtgatttcga gattcagttg 540
gagatattac acaatgcatt gcattcttta cttgggtggca aaggtaaata ttccatgtca 600
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atctgggcaa tctggcaaga ccttcagagg ttccgaaaac ggccataccg agaagcgaat 720
tgcgctatcc aattgatgca cacgccactc cagccgtttg ataagagcga caacaatgac 780
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gatttcgatc tggaaattga gattacggct gtgaatggat ctcatctaga cagtcatgtc 1200
atccactctc ccactatact gtttgaggcc ggaaca 1236

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<210> 99

<211> 1257

<212> DNA

<213> Megathura crenulata

<400> 99

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caattggaca agcgtcaaca actgtcactg gtgaaagccc tcgagtccat gaaagccgac 120
cattcatctg atgggttcca ggcaatcgct tccttccatg ctcttccctc tctttgtcca 180
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tcaactattc atgacccgga gacaggcaga gatataccaa atccatttat tggttctaaa 420
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cagggatcaa caaaaacaca tcataactgg tttattgagc aagcactgct tgctcttgaa 540
caaaccaact actgcgactt cgagggttcag tttgaaatta tgcataatgg tgttcatacc 600
tgggttggag gcaaggagcc ctatggaatt ggccatctgc attatgcttc ctatgatcca 660
cttttctaca tccatcactc ccaaactgat cgtatttggg ctatatggca atcgttgacg 720
cgtttcagag gactttcttg atctgaggct aactgtgctg taaatctcat gaaaactcct 780
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actgtactcg atagtggcct tattcccaca ccgtcaatca tctatgatcc tgctcat 1257

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<210> 100

<211> 1254

<212> DNA

<213> Megathura crenulata

<400> 100

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catgatatta gttcgcacca cctgtcgctc aacaaggttc gtcatgatct gactacactg 60
agtgaagcag atattggaag ccttaaatat gctttgagca gcttgacagg agatacctca 120
gcagatgggt ttgctgccat tgcattcttc catggtctgc ctgccaatg taatgacagc 180
cacaataacg aggtggcatg cagatcccat ggaatgccta cattcccca ctggcacaga 240
ctctacaccc tccaatttga gcaagctcta agaagacatg gctctagtgt agcagtaccc 300
tactgggact ggacaaagcc aatacataat attccacatc tggtcacaga caaagaatac 360

```

85

```

tacgatgtct ggagaaataa agtaatgcca aatccatttg cccgagggtg tgtcccctca 420
cacgatacat acacggtaag agacgtccaa gaaggcctgt tccacctgac atcaacgggt 480
gaacactcag cgcttctgaa tcaagctctt ttggcgctgg aacagcacga ctactgcgat 540
tttgagctcc agtttgaagt catgcacaac acaatccatt acctagtggg aggacctcaa 600
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```

<210> 101

<211> 510

<212> DNA

<213> Megathura crenulata

<400> 101

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aacacactta ccaactgcaga ggtggacaat ctcaaagatg ccatgagagc cgtcatggca 120
gaccacggtc caaatggata ccaggctata gcagcgttcc atggaaaccc accaatgtgc 180
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gaagaggaca cctttcca tcatggtcac atagactatt tgggagtggg tacaactcgg 420
tcgccccgag caaagtgtgt caatgatcca gagcgaggat cagaatcgtt cttctacagg 480
caggttctct tggcttttga gcagacagat 510

```

<210> 102

<211> 942

<212> DNA

<213> Megathura crenulata

<400> 102

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ggcctgccct actgggattg gaccatgcca atgagtcatt tgccagaact ggctacaagt 60
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gtggcgcttg aaaatggtgt aacaagcagg aatcctgatg ccaaactttt tatgaaacca 180
acttacggag accacactta cctcttcgac agcatgatct acgcatttga gcaggaagac 240
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tggggtcttg atagaatgta caagtatgag atcactgagg ctctgaagac gctgaatgtt 840
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```

<210> 103
 <211> 1248
 <212> DNA
 <213> Megathura crenulata

<400> 103
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 gtcgaagaac aaacttcgtt gaggcgagct atggcagatc tacaggacga caaaacatca 120
 ggggggtttcc agcagattgc agcattccac ggagaaccaa aatgggtgtcc aagccccgaa 180
 gcggagaaaa aatttgcatt ctgtgttcat ggaatggctg ttttccctca ctggcacaga 240
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 aatgatgcta tttccagcca ggaagaagat aacccatggc atcatggtca catagactct 420
 gttgggcatg atactacaag agatgtgcgt gatgatcttt atcaatctcc tggtttcggt 480
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 catgttggag ataatttctt ccttaaatat gaagcctttg atctgaatgg cgggaagtctt 1200
 ggtggaagta tcttttctca gccttcggtg attttcgagc cagctgca 1248

<210> 104
 <211> 1257
 <212> DNA
 <213> Megathura crenulata

<400> 104
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 aatatccggg acctctcaga gggagaaatt gagagcatca gatctgcttt cctccaaatt 120
 caaaaagagg gtatatatga aaacattgca aagttccatg gaaaaccagg actttgtgaa 180
 catgatggac atcctgttgc ttgttgtgtc catggcatgc ccacctttcc cactggcac 240
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 catggattcg acatcaaagt tgacgtcaga gctgtcaatg gatcgcatct tgatcaacac 1200
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 <213> Megathura crenulata

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 caagctattg cttctttcca cgctttgcct cctctttgtc caagtccatc tgctgcacat 180
 agacacgctt gttgcctcca tggatggct accttccctc agtggcacag actctacaca 240
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 acccattcaa tagtcatct tcattacgca tcatacgatc cactgttcta tatccaccat 660
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 gacaggcttt acaagtacga cattacaaaa acgttgaaag acatgaaact gcgatacgat 1140
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 <212> DNA
 <213> Megathura crenulata

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 gtgaacgggt atcaagccat tgcattcttc cacggtctcc cggtctcttg tcatgatgat 180
 gaggacatg agattgcctg ttgtatccac ggaatgccag tattcccaca ctggcacagg 240
 ctttacaccc tgcaaatgga catggctctg ttatctcacg gatctgctgt tgctattcca 300
 tactgggact ggaccaaacc tatcagcaaa ctgcctgatc tcttcaccag ccctgaatat 360
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<210> 108
 <211> 309
 <212> DNA
 <213> Megathura crenulata

<400> 108
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 aatggatatg aatcaatagc cggttacat ggtatccat tcctctgccc tgaacatggt 180
 gaagaccagt acgcatgctg tgtccacgga atgcctgtat ttccacattg gcacagactt 240
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